

| gtgcagcaac | tccagctttt | cttcattggc | taagtgccta | tgttttaaca | aaattta | 357 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| <210> <211> <212> <213> | 19114 436 DNA Glycine max | K | | | | |
| <223> <400> | unsure at a | all n locati | ions | | | |
| agctttttcc | anaactttgt | ttatcctgca | caaagcaatc | ttgtttggat | catgtctaca | 60 |
| gctttgaact | tattcccttt | tcatgatccc | catgttaggt | tattaactgt | gcttctagac | 120 |
| aggcatgttg | aaagaatttc | tcggagaatg | gtatgtctga | ctatgctaca | gtttcttaat | 180 |
| tagagggaca | ttctttagca | tcaatggatg | aaaaactaat | tgttgatact | ttgtgcagtg | 240 |
| caaccttcca | tatgttacta | tggtagtgac | tgacaatcta | caggttattc | tcacttcagt | 300 |
| tctattcaat | cgctcagtgt | ctaacatagt | tcgataggac | atgttgactn | tgtagccata | 360 |
| ttgcagtttc | tgcgttataa | aattgttntt | gggctctaac | ttagttgtta | gtgtttaaaa | 420 |
| tttatttttg | caatta | | | | | 436 |
| <210> <211> <212> <213> | 19115 311 DNA Glycine max | × | | | | |
| <400> | 19115 | | ٠. | | | |
| tcttatccaa | agcaattctt | ggtgttgaag | ctccttcttt | cttggcttat | tccctaatgg | 60 |
| atggtgcctc | ccctctcctc | ttctcctttg | ccttccgccg | catctccatg | gtgtaaaatc | 120 |
| accattgaaa | gacctcattg | aagctcaaaa | atccagcctt | catggaagct | ccacaagcaa | 180 |
| gcttccatca | agtggtaatc | aaagcacaag | agcttcaagt | aggtgctcct | taaacctccc | 240 |
| attaatggtt | tgctttaccc | tttcttgcat | tggtggttct | tcatttttct | ccatgtatct | 300 |
| cctcacatgt | c . | | | | | 311 |
| <210> <211> <212> <213> | 19116 405 DNA Glycine max | ς. | | | | |
| ~223× | uncuro at a | all n logati | ione | | • | |

| agcttgttag | ttgtaaattn | tgtaagactt | aattcacccc | cccccctc | ttaagttatt | 60 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| gaagccactt | gtccaacaaa | gggggatgga | tcccaaggtg | tgttcaaagt | atgtaaagga | 120 |
| attacaaaga | taatggaaaa | tctcaagtgg | attgtttgag | gactggacgt | atgcatggga | 180 |
| agtggccgaa | ccagtataaa | tcgagtgtga | aattctctct | tcccttattt | atntatttta | 240 |
| ttgcaatcaa | ttgtgtcttg | cacgtttaaa | gaacattatt | aaatcgattg | atgcttcttc | 300 |
| ttcttcattc | taagtctatc | atttaaaaga | aggttaacag | cttgttagtg | agaaattatg | 360 |
| tgagacttaa | ttcaccctcc | ctcttaagtt | attgagacca | cttgt | | 405 |
| <210> <211> <212> <213> | 19117 126 DNA Glycine max | × | | | | |
| gagagacatt | tgggagacgc | tgcttaccct | gccttcagcc | ttagactttc | tgggcctgaa | 60 |
| cgatgttatt | ggacaaaggg | atgaactccg | atacttttct | tatatgggac | atgatatatg | 120 |
| ctaaaa | | | | | | 126 |
| <210> <211> <212> <213> | 19118 222 DNA Glycine mas | x | | | | |
| <223> <400> | unsure at a | all n locat. | ions | | | |
| agctnttcgt | cttacagaca | gcaaaaagtt | tatacggata | accactcggg | tatttccgcc | 60 |
| cgtcagcgtg | actcaaaagt | caatatgaca | gatcttgtga | gcgcggaaga | tgacgtaaat | 120 |
| ctccgcgtgt | caacgggctt | gtcggccgcg | attgatgaat | ggcgcagaag | acgacgttag | 180 |
| tctctgcgtg | ctatcaggct | tttcgtctta | cagccacaaa | aa | | 222 |
| <210> <211> <212> <213> | 19119 442 DNA Glycine ma: | × | | | | |

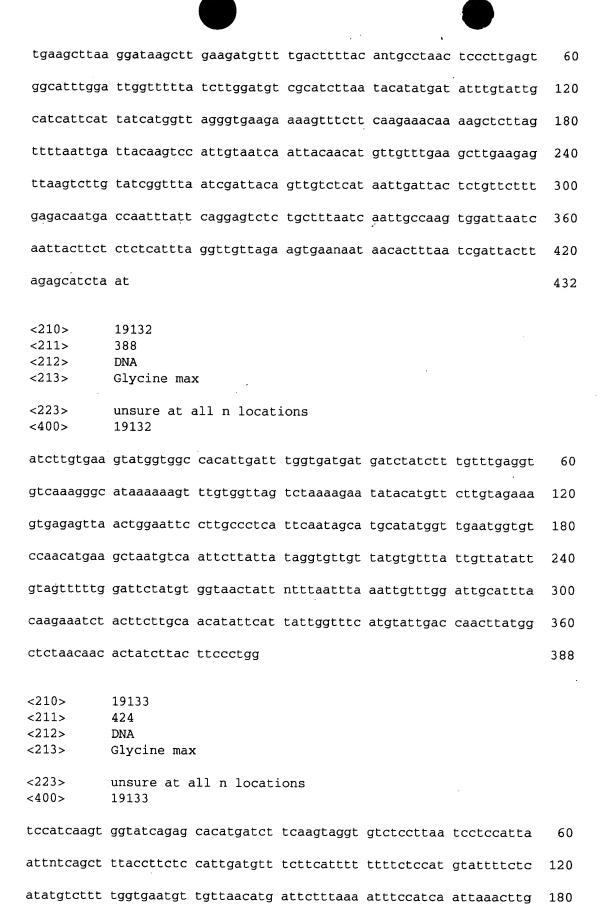
| acacgtggag | atttacgtca | tcttccgcgc | tcacattatc | tggcatattg | tcttttgagt | 60 |
|---------------------------|------------------------------------|-------------|------------|------------|------------|-----|
| cacgctgacg | ggcggaaata | cccgagtggt | tatccgtata | aactttttgc | tgtctgtaag | 120 |
| acgtaaagcc | ttataacacg | cagagactaa | cgtcgtcttc | tacgaccttc | gtcaatcgcg | 180 |
| gccgacaagc | ccatttaaaa | gcggagattt | acgtcatctt | tcgtgctcac | aagatctgtc | 240 |
| atactgactt | ttgagtcacg | ctgacgggcg | gaaatacccg | agtggttatc | cgtataaact | 300 |
| ttttgcattc | tgtaagatga | aaagcgtgat | agcacgcaga | gactaacgtc | gtcttctgcg | 360 |
| cccttcgtca | atcgcgatcg | acaagcccgg | tggcacgcgg | agaattacgt | catcttccgc | 420 |
| gctcacaaga | tctgtcatac | tg | | | | 442 |
| <210> <211> <212> <213> | 19120 324 DNA Glycine max | s. | | | | |
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| agcttttact | ttaatataag | tcctttattc | taaggttcat | aacacaaatt | aaagtgcaaa | 60 |
| gttgggatta | cttacttgga | ttgcaataag | acatgatcaa | gaggaagtca | tacaagcaga | 120 |
| gagcataatg | tgagcatagt | gcaaataaat | gcaagatgca | aaggatgata | gtgagaccat | 180 |
| gtttgtaaat | gcacgacaga | ctactgccta | aagcaattaa | gccttatttt | tggtagtttt | 240 |
| gactgcctcg | cttagcgcaa | gtcactcgct | tagcgagcac | tcaaggactt | ttaagttttc | 300 |
| agaatacaca | ctcacgagct | cagc | | | | 324 |
| <210><211><211><212><213> | 19121 434 DNA Glycine max | : | | | | |
| <223> <400> | unsure at a 19121 | 11 n locati | ons. | | | |
| ttatcactgg | cattgcactt | cctgttggtt | ccgacaaact | ttattctggc | atcactgatg | 60 |
| ggacagttag | gatatgggac | tgccatactg | gtcaatgtgc | taaagtcatc | aatcttggtg | 120 |
| ctgaaggtac | ctctttgatc | agtgaggggt | catggatttt | tgttggtctg | caaaatgctg | 180 |
| tcaaggtaag | ctcttatctg | gcattggttt | ggtttgatgt | atgataatgt | ctaatcataa | 240 |

| anather. | + aaaaaa = - | ++ =+ =+ == == = | ataattaat- | tassacete | gaatateese | 300 |
|---------------------------|------------------------------------|------------------|------------|------------|--------------------|-----|
| gagtagtaca | tgcaaactga | ttatgtggct | gcggctggtg | rgaaagcttg | yaatatccag | 300 |
| accatgtcag | aagttactct | cgatggaccc | aaaggccgaa | tccctgccat | gacttgtggc | 360 |
| aacaatacac | tctnttctgg | cgcagaggta | actaaccatg | ttattaatat | tgcgcaatga | 420 |
| tattccccta | accg | | | | | 434 |
| <210><211><211><212><213> | 19122 421 DNA Glycine max | ς . | | | | |
| <400> | 19122 | | | | | |
| agcttgttgg | ttacagtgac | aacaattggg | ctggagatga | agatgattgg | aaaagtacca | 60 |
| gtggatttgt | gtttttcata | ggaaacacaa | ccttcacttg | gatgtcaaaa | aagtagccga | 120 |
| tattcactct | tttgactcgt | gaggcagaat | acgtagcagc | tacttcatgt | gtttgtcatg | 180 |
| caatctagca | taagaattta | ttaaaagagt | tgggcatgtc | acaagaagag | ttgaccaaga | 240 |
| tctttgtgga | taataagtta | gtcattgctc | tagcaaggaa | tccagtgttc | tatgatcga a | 300 |
| gcaagcatat | tgatacccct | taccactaca | taagggagtg | catagcaaga | aaggatgta c | 360 |
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| a | | | | | | 421 |
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| tactcatttt | tggtgtgtga | gtttctggag | atgggcgacg | tcaagaagat | tttgaaggat | 120 |
| gatgaacaag | caattgcgtt | tgattggaat | aaaagggtgg | atgttgttaa | aggtgtaac a | 180 |
| aatgctttat | gctatatgca | tcatgattgc | tcacctccaa | tcgttcatcg | tgatatat ca | 240 |
| agcaagaatg | ttcttttgga | ttccgattat | gtagctcatg | tctcagactt | cggaacagcc | 300 |
| aagtttctta | atccagattc | atccaattgg | acctcctttg | caggaacctt | tggatatgct | 360 |
| actccaaatt | aatttccttt | ctctatacta | tttgagtaaa | tcatgatatt | ntagtttgtc | 420 |

| ttcgttagcc | atttacaaat | atatat | | | | 446 |
|----------------------------------|------------------------------------|--------------|------------|-------------|------------|-----|
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| <223> <400> | unsure at 19124 | all n locat | ions | | | |
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| cacctaaagc | cgagttcaat | ggtggttcgt | gccttcgacg | gcacccgccg | agaggttatg | 180 |
| ggagagatcg | atctcccagt | acagataggc | cctcacacct | gtcaagttac | cttccagata | 240 |
| atggatatta | accccccta | cagctgtctg | ttggggcgtc | cgtggatcca | ctcagtgaga | 300 |
| gttgttccct | ctacactcca | ccataatgtg | aaattcttag | tggaagggca | tctggtcatt | 360 |
| <210> <211> <212> <213> | 19125 443 DNA Glycine max | × | | , | | |
| <223> <400> | unsure at a | all n locat: | ions | | | |
| tctaagaata | gccttgataa | ctntaacata | atccattgat | tcttccccaa | aaaatatagt | 60 |
| gtcatcagca | aattgaagga | tattcactgc | aaccttgttc | ttccccacca | taaagttgtg | 120 |
| gaagcagtgt | ttggatattg | cttccttcat | cattcctgtc | aaacgttcaa | caaccaagtc | 180 |
| aaacaataaa | ggggccaaag | gatccccttg | tctcaaacct | ctttgaggct | taaattcagt | 240 |
| agttgggctt | tcattaacta | cgatagatat | agaggctgat | gtgaggcacc | ccttgaccca | 300 |
| actaatccac | ctgtcatgaa | accccattct | tctcatcata | taaaaaagga | atttccaaga | 360 |
| cacatagtca | tangctnttt | cgaaatccac | tttaaacacc | aagcaagacc | tctttgacct | 420 |
| cctaagcccc | tcaacaacct | cat | | .* : | | 443 |
| <210> | 19126 | | | • | • | |
| <211> | 443 | | | | | |
| <212> | DNA | | | | | |
| • | Glycine max | 7 | | | | |

| <223> <400> | unsure at 19126 | all n locat | ions | | | |
|--|--|-------------------|------------|------------|------------|-----|
| cttttttcc | tcatatnaga | acatacctag | ctaccatatg | tcggaattgg | accagtttaa | 60 |
| cagccacatc | taaaccggat | gacttctaat | ttcgaaaata | actatttcta | acttttccag | 120 |
| gggntaatga | actccaataa | cctcattcaa | aaccctgaac | tatctccact | ttcatcaata | 180 |
| catatgtcat | caagcagcaa | gtaatcccta | caaaagccat | atattgggta | gtacgctaca | 240 |
| taaatagtgt | cgaatccagg | cccatgttca | gcttcataca | agagaactga | gagtctctgt | 300 |
| cgataattac | ttggtagcac | cgatggtctc | aattcttttc | tctgactatg | acacgctgct | 360 |
| ctcatttgac | cacactctac | agaagataga | tgaatatatt | gaccacacat | ataanggaat | 420 |
| ngntggaata | gaannagcaa | cat | | • | | 443 |
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| atgcatatct | ggtaggtgaa | catcttaata | atgtggtgtg | ggatttagaa | cgcttt | 176 |
| <210> <211> <212> <213> <223> <400> | 19128 331 DNA Glycine max unsure at a 19128 | k all n locati | ions | | | |
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| gatgaggcca | aagaggtatc | ttttataaag | aaactagtta | ttcatttcag | tttacacaaa | 120 |
| aaaattaact | attttttcct | gttcaaagca | taattntgta | tttttttct | taagttatac | 180 |
| tgagttccta | aattatgttt | atatgtaagg | cccttgaaga | agtactttct | caaaattctt | 240 |
| caaagcagta | tgactcatac | tttgctaata | cacgtgaatc | tgttntccag | gtaataatgt | 300 |
| aatatactta | aaagaaccat | gtcatccttt | С | | | 331 |

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|--|---|---------------------------------|
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| actaatgaat | ctttgtgcaa ggagttttct attgacatgc caaatgagtt tgagatgtcc | 120 |
| atgatgggtg | agttaaacta ctttcttata ttacaaatca aaccaacaaa tgatgggatc | 180 |
| tttgtcaacc | cagcaaaata ttacaaggaa ctcatcatga aattcggaat gaagaactca | 240 |
| aaacacttgg | ctactcctat gagcactggt tgctaccttt gacaagatga atccggtcaa | 300 |
| ttcgttgatg | aaaagcaata tagaggtatg attggatctc tactttactt | 355 |
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| <223> <400> | unsure at all n locations 19130 | |
| atcttattaa | · | |
| | actttaaaca gagttaatta ttctcaaacc aaaatgaatt ctagcagatt | 60 |
| gttetggagg | actitaaaca gagttaatta ttctcaaacc aaaatgaatt ctagcagatt acctanggat aattctaaca gattgttctt ctggttttaa ctgtgaggtc | 60 120 |
| • | | |
| aatgttctcc | acctanggat aattctaaca gattgttctt ctggttttaa ctgtgaggtc | 120 |
| aatgttctcc gaaaaattct | acctanggat aattctaaca gattgttctt ctggttttaa ctgtgaggtc catacataat atttagaggc tataagatgt taaattttat tggaccaatt | 120 180 |
| aatgttctcc gaaaaattct tgaggtcaaa | acctanggat aattctaaca gattgttctt ctggttttaa ctgtgaggtc catacataat atttagaggc tataagatgt taaattttat tggaccaatt tggcatgctt ggttagggat aattttagca tattgttctg gttttaagtg | 120 180 240 |
| aatgttctcc gaaaaattct tgaggtcaaa tcaatacctt | acctanggat aattctaaca gattgttctt ctggttttaa ctgtgaggtc catacataat atttagaggc tataagatgt taaattttat tggaccaatt tggcatgctt ggttagggat aattttagca tattgttctg gttttaagtg gttccctata gcatattcta acatatctt gatacattg gctttgtcta | 120 180 240 300 |
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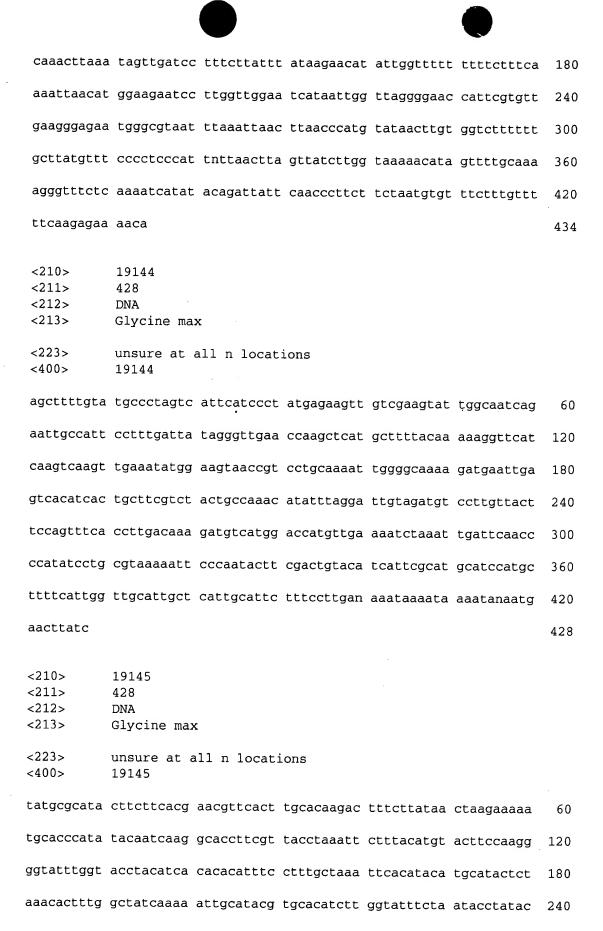
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| | | | • | | | |
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| aaccatgaat | tgtgttgagt | ttaggttcct | ttgagttttg | tcttgatatt | tttgtggccg | 300 |
| aaacctaaac | cataaaattc | ttactaaaac | atcaaagtag | atgaaaacct | caaaaatcta | 360 |
| gagtgatatg | ttcacttcat | tgagttttgc | ataaaagtca | tgctagtcat | gaaacttgca | 420 |
| cata | | | | | | 424 |
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| tcggcttaga | ttttcttcac | ggaaacgatt | tttccaagca | aattcgaaag | agagaggagt | 180 |
| gcaaaagggg | ctgaaccctt | ttcttcttcc | cttcctcccc | tatttatagc | aaaatatggg | 240 |
| aggtggttgc | cgcccagctc | gcccaggcga | gctcagctcg | cccaggcgag | ccaggttgct | 300 |
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| tgacaaccat | cgttttagga | gtgctgagca | ccagcagcgc | ttcgaggcca | ttaaaggatg | 120 |
| gtcatttctc | ccggagcgac | gcgtccatat | cagggacgac | gaatataccg | acttccagga | 180 |
| ggagatagtt | cgcccgcggt | gggcatcgct | ggttaccccc | atggccaaat | tcgacccaga | 240 |
| cataatcctt | gagttttatg | ccaatgcttg | gcctacagtg | gagggtgtat | gagatatgcc | 300 |
| atcctgtgtg | agggggttag | tggattccat | tcgatgcgga | tgctctcagc | cagttcttgg | 360 |

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|-------------------------------------|---|------|
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| agatttggtg | g agtcgatcta ctatgaccca ca | 92 |
| <210> <211> <212> <213> | 19137 344 DNA Glycine max unsure at all n locations | |
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| acacatgaac | gaaaacacaa ttcatggggc ttcgaaaaag gggttgagaa tggagaatta | 120 |
| cactaagcaa | tcactacgca tagctccaaa ctcgaaggtg gaggacacat gaacgataac | 180 |
| gcaattcatg | gggctccgaa aagattgaga atggaaaatt gcactacgca atcactacgc | 240 |
| atagcttcaa | acgcgaatgt ggaagacaca tgaatgaaaa cccaattcat ggggctccca | 300 |
| anagattgag | aatggagaat tgcactaagc aatcactacg cata | 344 |
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| gcgccatcca | ctatggaata agcctgcaag aaagagcttc accaccaaga tgagccttgg | 180 |
| ataagaagct | cggagaggat gcttcaatgg agaaaaagaa agagggagag aaagagagag | 240 |
| gtgggagcac | gaaattgaag gaagaaaaat aagggagaga agttgaactt tgagttgtgt | 300 |
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| ct | | 362 |
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| <210> <211> <212> <213> | 19139 313 DNA Glycine max | |
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| cgtgaacctc | acttgaggtg ggcaacaggg gatggggggc ttatgcgcgc tttgtgga | tg 120 |
| tggaaaactt | ggtgtgcacc aatcgccgac cgccacctag taccacatgt gatggata | cc 180 |
| ccataatcct | acaagcctga gatgaggaag tgtagaaggg tgaacttcct gcttttat | tc 240 |
| gttgaccaca | gagtggtacc tggagatatg tcgcgggggt caagagacct tggggacg | tc 300 |
| aggcggggtg | cta | 313 |
| <210> <211> <212> <213> | 19140 430 DNA Glycine max | |
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| tggaggaatc | ttctggaggg cccaagtggg cctggttgct atttgcaccc ccattttta | ac 120 |
| taaatacacc | ccttgccttt ttttggtgat tctttntttc gtaaagttac gaaaattta | ac 180 |
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| ccttttgact | tccggtgtgt cacagaacct tacnggatgt gcatcaatac tttcttttg | ga 360 |
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| gaccaaacac | | 430 |
| <213> | 19141 454 DNA Glycine max unsure at all n locations | |
| \UUJ/ | ampare at all infocations | |

| <400> | 19141 | | | | | |
|-------------------------------|------------------------------------|-------------------|------------|------------|------------|---------|
| tctattatct | taagttagag | gtgttcgtca | aaatgaanat | tgtttagcca | aagaataaaa | 60 |
| caaagaacaa | aaaccacatg | gaaagaaagc | aaaagaaaaa | aaacaagaaa | aagagagaga | 120 |
| gagagagaga | aagaatcaat | ccatccaaga | tggaagaaga | gagaaggaaa | atagaaaaga | 180 |
| aaaacaagtt | ctttggacca | gacaatgtct | aaaaaatgtg | cagaattgtc | ggaaagaaaa | 240 |
| aaataaaaga | gaagagcaat | agttatcaca | tgctttagtt | acaaaccaaa | tctttgtgtc | 300 |
| tgccctcctg | ttccacacca | aacaaaagag | aaagggaaac | agaaagagaa | aaggccgaaa | 360 |
| caaccaaagc | caaatttcct | accaaaatcc | aaccttataa | agacctattg | atccatgatg | 420 |
| attatgcata | ttatctttga | tttgatggga | aatg | | | 454 |
| <210> <211> <212> <213> | 19142 367 DNA Glycine max | | | | | |
| <223> <400> | unsure at a 19142 | all n locati | ions | | | |
| agcttcttcc | ttctttcgtg | ctctgntcct | tccttatttc | tggaggtgct | agacctcgaa | 60 |
| ttctagcctt | gatacattgc | tctaacattt | tgttttttaa | atttttatgt | tgatttgcta | 120 |
| acagattcga | tatttagatt | tttatgttga | tttgccatgg | atntgggttc | tcttgttctc | 180 |
| cttgattntg | gatttgtaac | tcctaaatct | gagtaaatgt | taagttgttt | ttgttgtcta | 240 |
| aatatgagat | ttgagttntt | tttttttt | tttttgtgtt | tctagtgtgc | tttntatgtt | 300 |
| gtatntatgc | ttttgttggt | gatgaacaaa | gaggggacgg | cgggttggtg | gtgtaaaaga | 360 |
| tggagag | | | | | | 367 |
| <210> <211> <212> <213> <223> | | k all n locati | .ons | | | |
| <400> | 19143 | | | | | |
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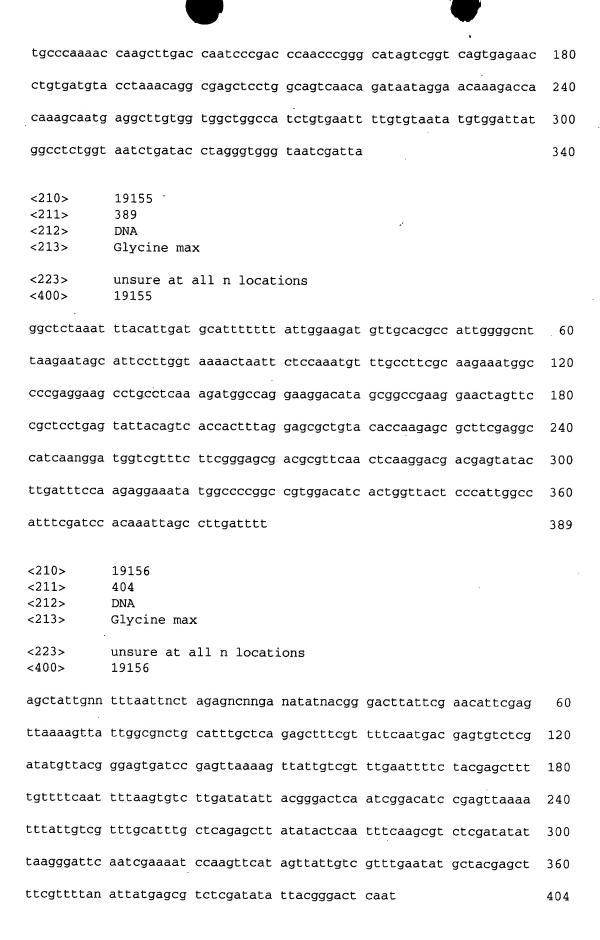


| atacacaaac | ttcatgatga | atcttgacta | tcgacacaat | aaggtgctac | atttcatgct | 300 |
|----------------------------------|------------------------------------|------------|------------|------------|------------|------|
| ctcctttttt | tcaagtattt | ttactaccta | aagccgcatg | caaattcaag | tatattntct | 360 |
| tttgctcact | aaaattgtat | tcaaattaaa | aggtatttt | gtaatgtatt | ttctgcaaca | 420 |
| tatttata | | | | | | 428 |
| <210> <211> <212> <213> | 19146 437 DNA Glycine man | x | | | | |
| <400> | 19146 | | | | | |
| agcttttact | atcctattca | aatgttaaca | tgactgttac | cctaaaataa | aatcaccaaa | . 60 |
| caaaagattg | ccaaaagtat | ctcccaccaa | ccccgaagat | caaatctcat | actccctccg | 120 |
| tttcaaaata | catgtccatt | tttgaaaaat | tgcggtaacc | aaggacaggc | taatttgaca | 180 |
| caaaagttcc | tattttaccc | ttgtccttta | ttttctccat | tttatattta | tttatcccac | 240 |
| ctcataatta | ctcccaatac | caaaattaat | taaagttaat | caaattacaa | taccaataca | 300 |
| tactggcaat | accaatacta | ctaaatggca | ttatgtttgc | ttcggtattg | aaaagctcaa | 360 |
| tgggcatagt | tcggttatca | aaagttttta | aaactcaatt | gaaaaaactt | ccctccatta | 420 |
| ttacatatac | tctaatc | | | | | 437 |
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| <400> | 19147 | | | | | |
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| cacaccttcc | acatgtaaac | tcaaccaatt | tcctctttaa | cttatgtcct | gtgacattgt | 120 |
| cctcatctac | agatettett | ctatttttct | ttggccttcc | tctttggacc | tttttatgtg | 180 |
| gtggaacaag | gtgtgtatac | tgtgtctggg | cccaatattg | aggtccttgg | actggttcaa | 240 |
| taaaatgctg | gtatgtctta | ttataagctt | ctattgacag | ccactcatga | cacatgtcct | 300 |
| caagcttccc | tcctttgtga | gttattgttg | caatggcatg | tcggcatggc | atccctacat | 360 |
| caaagttgta | aaatcagcac | acatgtag | | | | 388 |

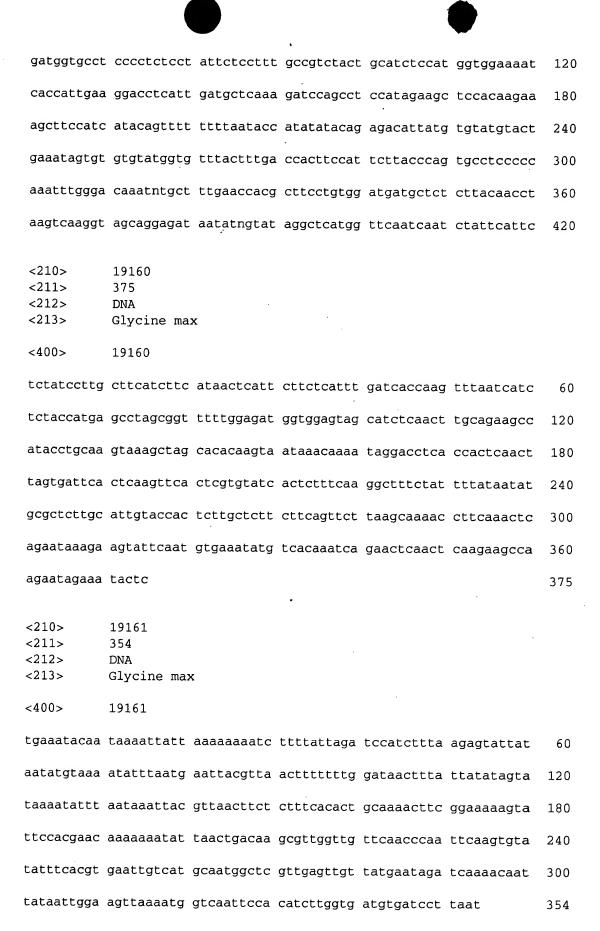
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| acaagctcaa | gtgcaacggc | agattccttt | gccttttctc | caacaacaga | aaggaggtat | 120 |
| ctactgtttg | gttttccttc | tccttgcatt | tgggaatggg | ctctatgcta | aactttccac | 180 |
| aacacttgct | cttgctcaat | tctctttact | gcactttta | gcatttcgtg | gcactatttg | 240 |
| tccaatcatg | gtttttacat | tccttctcta | actttttta | taatggctta | tcaattggag | 300 |
| aggaaagcaa | acatggataa | ccatgcattt | cctcttgttt | ggttgaagag | aaattgaaag | 360 |
| aaaagggaga | caaaatttgt | cttctagaaa | caaatttatt | ntttcctttc | attctcttat | 420 |
| caattcaa | | | | | | 428 |
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| tatgaccaaa | acaaaataaa | gaanaataaa | atgctacaat | gttttacagc | acaaggattt | 60 |
| gaaagggtaa | ccaccaagaa | tccccaatgg | actgtgattt | caatattatt | ttttgagaaa | 120 |
| aatatgtata | attaattttt | aaaatttaaa | tattataata | agtttatatt | taagcgaaag | 180 |
| taataagtat | aattataaaa | ttaatttgag | gaaattgaga | tttgaaaaaa | aaaattaatt | 240 |
| taaaaatctg | atacaaaatt | ataaatcatt | aattatttga | tagttntaat | aaatatataa | 300 |
| ttaataaaca | tttataaata | attaatgatt | ttgaaagttt | aataaatata | tataaacata | 360 |
| agtgctaaac | aatcataaag | ccatgtatca | caactcaaaa | agaaaatgag | tataataaaa | 420 |
| tatgtcttta | tattaaaaaa | caaataata | | | | 449 |
| <210> <211> <212> | 19150 509 DNA | | | | | |

| <213> | Glycine max | × | | | | |
|----------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
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| tactntagag | ccgacctcga | ggcatcgcat | ttcnatatgt | atatctatag | aagcctgagg | 120 |
| aactagcttc | agtttctacc | acacaattaa | tttaacaacc | ctatgggcca | agggtttcac | 180 |
| ttttcaacct | cataaagact | cggtctttta | tcttttctcc | cactagaaag | aagtaaacct | 240 |
| ataaatggaa | ataagcgctt | ggtttctaat | ttcttccaaa | acctttaaaa | ttttggaatg | 300 |
| atgtctgaga | caaaaccatn | tcagtattta | aagggcagcg | taaattgtgc | atgccaatat | 360 |
| cagtctcctt | acatccaagg | tcttacgaac | aacatacatg | ctgtataaag | ctatgggccc | 420 |
| ctgctgctca | cgggctgttg | catacataac | agcgaggaag | aggcccacaa | acgatggacg | 480 |
| ccaatagcac | cacaagatgt | gaccgcgcn | | | | 509 |
| <210> <211> <212> <213> | 19151 482 DNA Glycine max | ς | | | | |
| <223> <400> | unsure at a | all n locat: | ions | | | · |
| cgcagctgcc | tttgatgcgt | tgattgccac | nttgaaaacc | ggaccacaga | atctcagctg | 60 |
| accagcttga | tcggaaatca | tttttggcta | gttttataag | cntcggggat | gttgcccgta | 120 |
| atggtatttt | ccctggagcc | agcacttgcc | ctatagcatg | gaatctcttc | cagaagccgc | 180 |
| ttggtgttcc | taaaaaagct | tcctatatca | ctctgtgagt | taatttggta | ccggccagac | 240 |
| aaatattttc | ttcatataat | gtgccatttt | aatagttctc | tacttagttt | ctgcaggttt | 300 |
| acacgtgaac | aaggtgatga | caactatagc | tgtggcttta | ccactccata | tgaaccctgc | 360 |
| cttctggaga | gatgtgtaca | taaaatcaac | caccgccatg | cctcctgtgc | ctacacaatg | 420 |
| gaaggattat | gaagaattgt | gcctcccatt | ttgatggaca | ctgtcatcat | tcacgatatt | 480 |
| cn | | | | | | 482 |
| <210> <211> <212> | 19152 410 DNA | | | | | |

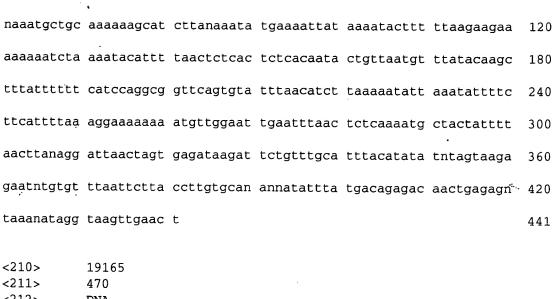
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| cagcttcttc | c tectacecag aaagggtaag ttatattgte taettteaca ee | cctccctgc | 120 |
| tctaactctt | t tgactccatc tttgtctata tggatgttac ttagagcagt at | gttttcat | 180 |
| ttgcgtgtgt | t ctatatatat gtgtaatgtt cagtgctgag tatgatattg ta | agațgttac | 240 |
| tgtacctcaa | a tgtgattett agatateata taatetttgt agtttgtage at | cttctgtt | 300 |
| aatattatct | t ctgcatttct tacctatatt tggattgtct tcttctttct aa | acttacat | 360 |
| ctggtcttat | t gtgctaatca ggtgaagtta catgacacag acattaacac | | 410 |
| <210> <211> <212> <213> | 19153 384 DNA Glycine max | | |
| <223> <400> | unsure at all n locations 19153 | | |
| ntaaacctct | cacaaaaggg aagactttta tagaagatgt atctccaatt to | cttagaggt | 60 |
| gagttcaaat | ccttacatat aaaagagtca gaatccattt tccattattt tt | taagattt | 120 |
| cttgttggtt | t cacatcaaat aaaaagaaat ggtgagaagt tagaagatgt ta | agaattatg | 180 |
| gagaaagata | a ctacgcccgt tagatcccaa atttgagcat attattgtga ca | atcaagga | 240 |
| aaccctagat | ttaaaaacca tgatgataga acaacttcaa ggatcattgc aa | igcttatga | 300 |
| agagaagcat | aagaagaagc aacagatcac taagccactc ttcaagatgc aa | ictgatgga | 360 |
| gaaggaagaa | a agtcaacgaa atga | | 384 |
| <210> <211> <212> <213> <223> <400> | 19154 340 DNA Glycine max unsure at all n locations 19154 | | |
| ttcttttatg | g agtaagtngc gaacggtgaa acttcctgct tttattgttg ac | cacagagt | 60 |
| ggtacctgta | a gatatgtegt gggggteaeg agaeettggg gaegteaggt gg | ggtgctat | 120 |



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| ctcagcttga | gcaagcaaca catactttta ctcgatgtct atcatccgta tattcagaca | 60 |
| ctagaaatga | aacgaactcg tacaattcaa atgacatact tttaactcga gtggattgag | 120 |
| tctcgaatat | atcagacete gtattgaaaa tggagetegt acaatgeaae gatataaett | 180 |
| ttactcgatg | tacgattagt ccgtatatat tgagacctca aattgataca aagcttgagc | 240 |
| aatgcaacca | catactttac tcgatatcga tgatccgaat aatcagagct cgaatgatac | 300 |
| gaactttaca | a | 311 |
| <210> <211> <212> <213> | 19158 398 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19158 | |
| ttcttattac | tcttccaatg gttttcttga ccagacatga agagtctata aaagcaagac | 60 |
| cttgacttgc | attcaaagaa cttcttgaac aactcttaag aaaccttgaa acctttacaa | 120 |
| cctttacaat | totttaagaa ttoattooca atoatettto ttottottoo tttgccaaaa | 180 |
| agctttctaa | gttttttgtt ttccaaacct tattcttctg caagtgaaaa ttctgcagaa | 240 |
| aacaaaagtg | tgctatatct tttcattctc ttcttccttt gccaaaaaga attcaacaag | 300 |
| gactaatcgc | ctgaattctn tntgtgtctc tcttctccct ttttccaaaa gtatagaggg | 360 |
| accaaccgcc | tgaattettt tgtgteteet tteteece | 398 |
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| tcttatccaa | ggcacattnt tggtggtgaa gctccttctt tcatggctta ttccctagtg | 60 |



| | <210> <211> <212> <213> | 19162 308 DNA Glycine ma | x | | | | |
|---|----------------------------------|------------------------------------|-------------|------------|------------|------------|-----|
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| | agcttatact | tctattttta | gcgtctcgag | cagttacggg | actcaatcag | acatccgagt | 60 |
| | taaaagctat | tgťcgtttga | atttgcacag | aggttcaaca | ttcaatgtcg | agcgtctcga | 120 |
| | tatgttacgg | ggctcactct | gactttcaag | taaaaagcta | atgtcgtttg | aatgttctca | 180 |
| | gagattctac | attcaattac | gagcgtctcg | atatgtgacg | ggactcaatc | agacatccga | 240 |
| | gaaaaacgtc | actgccgntc | gaattagctc | ataagttcaa | cattcaatgt | ctagcgtctc | 300 |
| | gatatatt | | | | | | 308 |
| | <210> <211> <212> <213> | 19163 416 DNA Glycine max | ζ | | | | |
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| | tctttataca | cctcgcagca | ctgagtgact | atcacgtagc | catgacctca | tagggacata | 120 |
| | agagcacaga | cagctgtagc | attcttctgg | acaagaccta | aggatttctc | ctgatatcgc | 180 |
| | cgacacgttc | ataactagca | ctaaaaccta | tgggggaagg | aaaaatagat | ctaatcttga | 240 |
| | cacaattaac | agacattaat | tgacccgtag | attgcactat | catagcaaat | atgcctacta | 300 |
| | ccgagactgg | cttgtgagga | cccccaatgc | gatcatggct | ttcaattata | cagtactgaa | 360 |
| | actggcccct | gggggaaatg | ctgacatata | ctgataacca | tttttcgaaa | gtcatc | 416 |
| | <210> <211> <212> <213> | 19164 441 DNA Glycine max | | | | | |
| | | unsure at a 19164 | ll n locati | ons | | | |
| , | agcnnttntt | aagaagtaat | cnnnnnnnca | tacatgtggn | ttagtgtatt | taacatctta | 60 |



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|-------------------------|------------------------------------|
| <223> <400> | unsure at all n locations 19165 |

| <210> | 19166 | |
|------------|---|-----|
| <211> | 394 | |
| <212> | DNA · | |
| <213> | Glycine max | |
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| agcttttgat | tattcattat tgattcaaag aatttttat gataacaaag gtgatgacaa | 60 |
| | and a substitution of the | · · |

agettttgat tatteattat tgatteaaag aattttttat gataacaaag gtgatgacaa 60
aaageteaaa ggteaagaac aetteatgat aacaaagatg atgeteteaa gaateaaaga 120
atgagtteaa gattgaatea agaacaette aaggtteaag aggaaatttg attteaagaa 180

| tcaagaatca | agtttcaaga | ttcaagttcc | gagaatcaag | atcaagattc | aagactcaag | 240 |
|--|---|--|--|--|--|--|
| attcaagaat | caagagaaga | cttaatcaag | ataagtatga | aaaagtgttt | tcaaaaactg | 300 |
| agtagcacat | ggattgttct | canaacttgt | ttaccacaga | agttntaatc | tctggtaatc | 360 |
| gattaccaga | ttgttgtagt | cgattaccag | tagc | | | 394 |
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| taagaataat | ggcccactgg | taaaactaac | tttccaaatg | tttgcct | | 107 |
| <210> <211> <212> <213> | 19168 427 DNA Glycine max | | | | | |
| <223> | unguro at a | .11 1 | • | | | |
| <400> | 19168 | ıll n locati | ions | | | |
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| ntggagtaa | a aggcttact | a aagaaggcaa | tagggtggcd | tittctgcgat | aagactgcac | 60 |
|------------|--------------|--------------|------------|--------------|------------|-----|
| ccatgtcga | c accaaacac | g tttgtttcca | aggaaaaagg | gataatgaag | tccggcaaga | 120 |
| gtaaagtag | g ggctatagaa | a agcacatcct | tcaatccagt | gaaaacttgt | tgcgccttag | 180 |
| atgaccact | c aaaggggtad | c ttcataagta | acctggttaa | ataagctgca | atggaggcat | 240 |
| agccacgaa | aaatcggtga | a tagaaacccg | agaggcccaa | aaaactgtgc | aaggccttgg | 300 |
| aggaatgag | g tgtgggtcad | tgctaaatgg | ctnggacctt | agccgacact | ggttcaactc | 360 |
| cttgtgcaaa | a aaccaggtga | cccaagaact | caacttgttg | ctgggcaaag | gtgcacttgg | 420 |
| atagt | | | | | | 425 |
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| acaattggcc | acataaattg | gattccaaag | gtttttgctg | tgtgaatagt | acctagacgc | 120 |
| ttaaacacaa | accttcttt | attcaagaaa | acataaaccg | tttcatgact | catttcaacc | 180 |
| attaatgggt | taaatgaata | attaatttct | agcaacaatc | aacagttttc | agagatgaat | 240 |
| agctgacaac | ccgtactctt | caactattag | tatgc | | | 275 |
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| <211> | 276 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | ĸ | | | | |
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| agggaggaac | ccatgttgtg | attgccattc | ctacatggcc | aaatttctca | acagctcaac | 120 |
| aatgtcaata | cttaaccaat | atcaaccctt | cttattaccc | accaccctat | cagccaaaaa | 180 |
| cacctaatca | tccacaaagg | ccacccctaa | atcagccaca | aagcccggct (| accacacatc | 240 |
| tgatagtaga | caccaccctt | aacatgaacc | aaaaca | | | 276 |
| <210> | 19172 | | | | | |

| <211> | 432 | | | | | |
|----------------|-------------|--------------|------------|------------|------------|-----|
| <212> | DNA | | | | | |
| <213> | Glycine ma | x | | • | | |
| <223> | | all n locat | ions | | | |
| <400> | 19172 | | | | | |
| agcttgtttc | aagatctaca | acaggaactt | gaaatgaaag | attcaatgag | agtgaaggag | 60 |
| ttacataatg | aaaattatga | ttcacagggt | acttgtgatc | attccttctg | tgataaggag | 120 |
| ctaaatggat | tntcacctga | aaagcacaca | gataactctc | caataactga | ctacaaaaaa | 180 |
| tcatatgatc | aaaaggaaga | agaaagatca | gaatctatga | gcaaaattga | agctgagctt | 240 |
| gaagctgaac | ttgagagatt | gggattaaac | atgaacgaat | ctagcccaga | aagaccgctg | 300 |
| tctgagcttg | ttgaggtaag | cataaaatgt | ntatcttttg | catttcctag | taatgacatg | 360 |
| atcctgtaag | canatttcat | tgtcttggaa | tatggctgct | gaaatatact | ntctttanat | 420 |
| aatntcttat | at | | | | | 432 |
| | | | | | | |
| <210> | 19173 | | | | | |
| <211> | 435 | | | | • | |
| <212> | DNA | | | | | |
| <213> | Glycine max | ζ. | | | | |
| <223> | ungung of a | .11 - 1 | | | | |
| <223> <400> | 19173 | all n locat: | lons | | | |
| 11007 | 171,73 | • | | | | |
| ctggactcta | gctagagaag | ctntaanaac | ctttattttc | atagttatat | atatatatat | 60 |
| atatatatat | atatatatat | atatatat | atatatat | atatatatat | atattttata | 120 |
| tgagaggggg | gaaaatatat | tatattaaat | cctaattctt | tatatatgaa | tctatgggct | 180 |
| agatataatt | ttctcatctt | atataagata | ttttctcgta | taagatatgt | acttatatat | 240 |
| gcacatattc | tttattttat | tgagaggtta | tttttaaatt | aatgagtttt | ttttttaaa | 300 |
| aaaaatctct | ataaaaacta | ttgggaaaca | aattaacata | atttcgtttg | cttttgtgtc | 360 |
| gcgttgatgt | caattctcct | tttataatta | tctattctac | tatgacggtt | cgcacgacat | 420 |
| aatttcaatc | ttaag | | | | | 435 |
| | | | | | | |
| | 19174 | | | | | |
| <211> | 437 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |

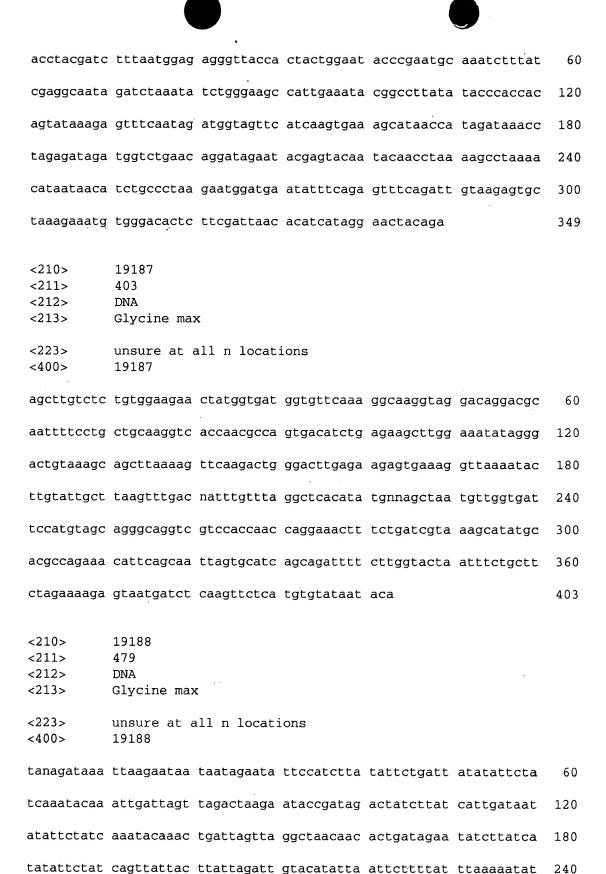
| <223> <400> | unsure at all n locations 19174 | |
|-------------------------------------|---|-----|
| ttctttgcgc | gtttttggag gctcagcgcg gatctggagc ttaacgcaca tttggcggct | 60 |
| tagcacacga | tcacttatgg cagcaaaact ttgcatgtcg cttagcgcgc agtgtaagct | 120 |
| tagcgtacaa | tcaatatcga aaaacataac tgtgctgtgg agaaaaaagg gagaaaccaa | 180 |
| aagaaagctt | ttttggaacc aaataaggag atagggcacg agagaagatg gagaacccac | 240 |
| tcaattgggg | accatttcct ccattttctt ccacacctct tgtttccttt ttgtattatt | 300 |
| aattntctca | tgacaatgag aggttaaacc attcactgtt ggaagctcaa caaccaaaca | 360 |
| ctctngatat | aatgatncta actatctatn taatgatatt ttgatattat ggggtctntt | 420 |
| ctatgctaaa | tatcatg | 437 |
| <210> <211> <212> <213> | 19175 398 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19175 | |
| aaaacacaat | tttgcgagat gtgttacaat tnttagtggt cttgatttcg agattaagta | 60 |
| tatcaaggac | acttcaaact ctctacttga ctatcttacc cgtgaattct tacagaaaaa | 120 |
| ttgccatgcc | acctaaggca tctagcacct ctttatgagg aggaagaagt tcaagtaaag | 180 |
| atttcaaatt | gactctacca gagccattct ccaaaaagaa ttcaccctca aaatctgggt | 240 |
| caccaaccca | agttgggtta ttaactcaga aaccaaaaca agaagggtca tctatccaac | 300 |
| tagtttcaat | taaacctaag tcatccactt aggaatttcc taaaaatcaa acattaaaac | 360 |
| ataccaaggt | cgactatgcc tttctaatag aaacacta | 398 |
| <210> <211> <212> <213> <223> <400> | 19176 372 DNA Glycine max unsure at all n locations 19176 | |
| tgcttttctc | ttagagatcc aggaaggata aagcggctga aggaaccagt tccgctctcg | 60 |
| aatatgacag | ccaccgtttt aggagcgctg agcaccaaca gtgcttcgag gccatcaagg | 120 |

| gatggttatt | tctccgggag | caacgcgtcc | : agctcaagga | a cgacgagtgt | atcgacttcc | 180 |
|---|---------------------------------------|-------------|--------------|--------------|--------------|-----|
| aggaggagat | agttegeega | tagtgggcat | cactagttac | ccccatggcc | : aagtntgacc | 240 |
| cagacatagt | cctcgaattt | tatgccaatg | cttggcctac | : taaggagggc | gtgcgagata | 300 |
| tgcgatcctg | ngtgangagt | cagtggatcc | cgtttgtgca | ı gatgctctca | gtcagctcct | 360 |
| gggatatcct | ct | | | | | 372 |
| <210> <211> <212> <213> | 19177 382 DNA Glycine ma | | | | , | |
| <223> <400> | unsure at 19177 | all n locat | ions | | | |
| tgctgcctca | tgaagaatgc | cttgctctta | gatagcatga | ttttagccct | tttataatat | 60 |
| ggatgtatgg | aaatatgtag | catgaaatgc | cttgcaaaat | gttgaatgaa | atgccttgcc | 120 |
| aaatgttgaa | taaaatgcct | tgcaaaatgg | tgaataaaat | gccttgccaa | atatgaatat | 180 |
| atatagcatg | aaaatgcctt | gcataatatg | aatatatata | gcatgaagtg | ccttacaaag | 240 |
| tggttgaatg | ggtagcgtan | aagtgttttt | aaaatatgtc | atttatgata | ggtggaaaag | 300 |
| aaccttccaa | aaaatgtgtg | tatatatata | ggatgtagca | tgaaaaggtt | tgtcaacaaa | 360 |
| atatatgtgt | acatggatgt | ct | | | | 382 |
| <210> <211> <212> <213> <223> | 19178 381 DNA Glycine max unsure at a | | .ons | | | |
| agctttgatt | nttatgtcaa | caaaaatagc | gattnttatc | acaataggtg | tgttttgttt | 60 |
| gtagttcttt | caaagcttaa | tttttttggt | taattgtttg | ggtgttcttg | atccttctcc | 120 |
| tactttgttt | cgcatagaag | cccgaagaac | acactaaagc | tttgattttt | atctcaagaa | 180 |
| aagcagcgat | ttttatcaca | acaagtgcgt | tttctttgta | gttctttcaa | agcttaattn | 240 |
| tttggttaa | ttgtttgggt | gttcctgatc | cttctcctac | attgtttcgc | atagaagccc | 300 |
| anagaacact | anagctttga | ttnttatctc | aagaaaaagc | agcgatttta | tcacaacagg | 360 |

| tgcgttntc | et tttgagttet t | 381 |
|----------------|--|--------|
| | | |
| <210> <211> | 19179 368 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <400> | 19179 | |
| tgtaatcga | t tacacacata ctgtaatcga ttaccagatg tatttttcag aaaacatto | ct 60 |
| caacagtca | c atctttttat ctgattctta agtggccatc aaaggcttat atatatgtg | ja 120 |
| ctagagacad | c gaattgaaca agagttttga agaacaaaaa ggtcttatcc tcttaacaa | ıg 180 |
| caaaattggt | t ttatcctctt acaaattcct tggccaaaac actcgtgatt caataagga | a 240 |
| ttatttgagt | t geteaaattg tteaatetat etettetag agagatttet tettetett | c 300 |
| ttctttattc | c tgaaaaggga ttaagagacc gagggtctct tgttgtgaaa ggattctaa | a 360 |
| cacaaatg | | 368 |
| | | |
| <210> | 19180 | |
| <211> | 306 | |
| <212> <213> | DNA | |
| <213> | Glycine max | |
| <400> | 19180 | |
| tttttgtaaa | a actttttgaa taaaatgtag cagaagttaa caataattct taaaaaaaaa | t 60 |
| gtgtggctag | cacttgttat tttagtatcc aagctaacct tcatatcgta cttaaatcca | a 120 |
| ttaacggcgt | tattgtctta catgccttta ttgtgacaaa ctattagact aataataata | a 180 |
| ttttaaaaca | tacttaaatg aattaataat ataaaaatat tctaatattt gtgacaaaat | 240 |
| aaatagattg | agagaattag agatactatg gaaatatttt acattgctac cactatttta | a 300 |
| catata | | 306 |
| <210> | 10101 | |
| <211> | 19181 299 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <400> | 19181 | |
| gcacgagaaa | acticaatcc acccaagaag gigitaaaaa accagagiic attccattat | 60 |

| cctagtttaa | gagaatctta | ccatgagaac | caaatctaaa | ctctgaagaa | gccaaacaag | 120 |
|-------------------------------------|---|-------------------|------------|------------|------------|-----|
| ataaaatcta | tgtctacata | cactgatggc | taaaccattc | caaactaaat | ggcaacttta | 180 |
| gcttaagatg | gctcataata | gttaaagctt | ggagaaaaag | ataacaatga | tgggagggac | 240 |
| agtatggtgc | ttatgatttt | acttttcact | gctctatgat | acataaccca | ctgttgagc | 299 |
| <210> <211> <212> <213> <223> <400> | 19182 377 DNA Glycine ma: unsure at a | x all n locat: | ions | | | |
| tagcttcttc | aaagacttgn | ggtccaacat | aggcatcatt | tcaagcctaa | tcaacgaggt | 60 |
| aacaccccct | tgggtggtcc | taacaattgg | tggggttctc | aattttttg | ggtatttcat | 120 |
| aatttggctt | gcagtggcca | gaaaaattgc | taagccccaa | gtttggaaca | tgtgcttgta | 180 |
| catcttcatt | ggagccaatt | ctcactgttc | caccaacact | ggagtcattg | tcaccagtgt | 240 |
| aaagaacttc | cctggcacaa | ggagcattgt | aattggcctc | ttgagtgggt | atcttggctt | 300 |
| gagtgcagct | atcatcactc | agatatacta | tgccttctat | ggaaatgatt | ccaagnttct | 360 |
| aattntgctc | atggcat | | | · | | 377 |
| <210><211><212><213> | 19183 375 DNA Glycine max | ς | | | · | |
| <400> | 19183 | | | | | |
| tctcgaggaa | gcctcttaat | gaagcttctc | tatgaagaat | tatgaagctg | ccttggtaaa | 60 |
| aatgcttccc | cacctttggt | aaccggtggc | tcttctcaaa | atttggtctg | gcgcttcaca | 120 |
| gaacacttgg | ccatgatctg | accggtggga | tctttaagaa | aatgtctgga | atgtgtgcca | 180 |
| tgttttcgct | tccgaaagca | ttgctcactt | gtgcggtttg | agccttgtag | tctaagtacc | 240 |
| tttggaaaaa | tgccatttct | ttccctttct | ttcttccaaa | accattttca | acattccaag | 300 |
| atctttctcc | atcacccaca | gccaccatta | gccaccacaa | actgtccgtg | ttcttcattg | 360 |
| aaaccccaca | ccgag | | | | | 375 |

| <210> <211> <212> <213> | 19184 383 DNA Glycine max | |
|----------------------------------|--|-----|
| <223> <400> | unsure at all n locations 19184 | |
| agcttgtacg | ctcatcgttc gcgtgtatga tatccactcc acacgatntg aagtagagga | 60 |
| gagcttcaac | cctataacgc aacgtggcgg acaaaagtgg gcagtaaact tgaatggccg | 120 |
| tcattgtcaa | tgcggaaagt attctgcgct tcactatcca tgttcacaca ttattgcagc | 180 |
| ttgtggttac | gtgagcatga accaatatat agatgttgtt tatacaaacg agcacatctt | 240 |
| aaaagcttac | tccgcacaat ggtggcctct tgggaatgaa gcggctattc ctccttctga | 300 |
| tgacgcatgg | acacttatcc ctgacccaac tacagttcgt gcgaaaggtc ggccaaaatc | 360 |
| aacaaggata | agaaatgaga tgg | 383 |
| <210> <211> <212> <213> | 19185 380 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19185 | |
| gagcgttaga | tgatattgtc ttcaccgacg aaaggatcaa agtgagtcta aaaagagaca | 60 |
| aatctgagca | tcatactttg ataaatgcca aagaaactat ggcaagtgaa gaggatgaga | 120 |
| aggagggaga | aacctatgtt gtgactgcca ttcttatacg accaagtttc ccaccaaccc | 180 |
| aacaatgtca | ttactcaacc aataacaacc cttctcatta cccaccaccc actcatccac | 240 |
| aaaggccatc | cctaaaatca accacaaagc ctacctaccg cacttccaat gacgaacacc | 300 |
| acctttagca | taaaccaaaa cacccaccaa gacatgaatn ttgcagtgaa aaagcctgga | 360 |
| gaattcaccc | ccaaatccag | 380 |
| <213> | 19186 349 DNA Glycine max | |
| | | |



cattgtcaaa gtaggataaa tatttaatta tatacaattn gttttattaa ttattaataa

| ccttctactt | tatcctaata | atttatactt | aacagatgag | tatttntctc | attataacta | 360 |
|----------------------------------|------------------------------------|------------|------------|------------|------------|-----|
| tgtactatcc | aatgtcttat | tagattttta | ataacatatt | atccatttaa | tttacagatg | 420 |
| actgtaattt | gatcaatcat | tagtatatat | atttatacta | tcgaccgatt | aagttaaag | 479 |
| <210> <211> <212> <213> | 19189 399 DNA Glycine ma | x | | | | |
| <400> | 19189 | | | | | |
| agcttctctt | gtaccttatg | caaatcctca | actcatcctt | caagatcaaa | ctgtctacta | 60 |
| gtgattggtc | cctttcctct | ctccggagct | taagctcgct | gttactgctc | cacagagccc | 120 |
| ctcggaattt | gttccggcca | tgtttttccc | tacgggccct | tttggtctct | tgttacaaag | 180 |
| ccttggtggt | ggcaatattt | acgtctcaga | gttcggcatt | ctcctttcgg | atcttaagag | 240 |
| ctgctgattt | gaacttgttt | ttgactgttt | ggactttctc | gagttctgcc | ttgagggctt | 300 |
| gcacctcttc | gtcctcgtcc | ggagcttcaa | cttccacccc | cttagtggtt | ctcaaactcg | 360 |
| ggagccaatc | cagaccttgc | atgtgggctt | tcaaccatc | | | 399 |
| <210> <211> <212> <213> | 19190 443 DNA Glycine max | ĸ | | | | |
| <223> <400> | unsure at all n locations 19190 | | | | | |
| ntaaggaagc | aatgcattac | cccacatgga | tcaatgcaat | gcgattatct | gctgaattct | 60 |
| attgagaaga | attcaacatg | ggaacttgtt | aatctgcctc | ttgacaagaa | acccatagca | 120 |
| ctgaagtggg | tttataaagt | gaaggtgaaa | tccaaatgag | gccagacttg | tggcaaaagg | 180 |
| gttcttatga | aaacctggag | ttgactatgg | tgaggtctat | gcacctgtgg | caagaataga | 240 |
| aacagtgaga | ttggtggtag | caattgcaaa | tatataaggt | tggtctatgc | ataaactaca | 300 |
| tgtgaagtct | gctttcttaa | atggacagct | agatgaggaa | gtttatgtgg | accagccact | 360 |
| tcttgagaca | ttgggacaag | atgaaaaggt | atacagattg | aaaaaggaat | atatggtctt | 420 |
| aataagctcc | atggcttgga | aca | | | | 443 |

| <211> <212> I | 19191 383 DNA Glycine max | |
|-------------------------------|---|-----|
| <400> | 19191 | |
| agcttgtagg g | gttcattcca tattccgttg tcatatgcta aacttgatcc catatccact | 60 |
| caataattca a | atggtagcca taaccccaac caaggttcct caacctccat ttttctgagg | 120 |
| atacgactcg a | agcgcaacgt gtgcttatca tggaggagcc ccgggggcatt ccattgagca | 180 |
| | ccgaagcata aagtgtgatg tctaattgat acgggctcgc tgaaattcga | 240 |
| | tgttgaatc ctaacattga caagcaacac catacatggg gcaattctgg | 300 |
| | atgactcat catgattatc aagtttatgc cataaaccac agttactatg | 360 |
| ctaaatgata t | ggataagat gga | 383 |
| <211> 4(<212> DI | 9192 65 NA lycine max | |
| | nsure at all n locations 9192 | |
| tctgtgagag ca | actteettg agaagetaga gettagetae acacaceett tteatateta | 60 |
| agctcacctc ct | ttgagaagc ttccttaaga agattcctaa agaagctaga gcttagttac | 120 |
| • | caatageta ageteacete ettgagatga gaagetagag ettagetaca | 180 |
| | atagctaag ctcaccccca tgacaaaata catgaaaata caaaaaatat | 240 |
| | gctactca taatgcctcg aaatacaagg ctaaaaccct atactactag | 300 |
| | acaaggcc ccaatgaagg anaaacctat tctaatattt acaaagataa | 360 |
| | ttagccca tgggctcgaa atctacccta aggctcatga gaaccctacg | 420 |
| ggccttcctt gg | atctctgg cccaatctac ttggactctt ctatc | 465 |
| <211> 377 <212> DNZ <213> Gly | A ycine max | |
| <223> uns <400> 191 | sure at all n locations 193 | |

| agcttttcat | gtctatggtt | tctagagaga | gaaaggtcca | agttctagag | agagagaaag | 60 |
|----------------------------------|------------------------------------|-------------------|------------|------------|------------|-----|
| gtccaagttc | tagagagaga | gaaaggtcca | agttctagag | agagaaaggt | ccaagttcta | 120 |
| gagagttttg | aaagattttg | ttgtgtgaag | atcagaagag | accaaagctt | gaaacaagag | 180 |
| ccggtttaag | agcttgagat | aagtttgtga | gtgattgtga | gatcctagag | gtgaaggaga | 240 |
| catcctcacc | acttgtatat | ttgcaatctt | tcatcttgtt | cttctctttg | ttcttaagaa | 300 |
| ggcttcctgg | tatggaactg | tggctcttcc | ctataggtac | ttgatgtaaa | tatatntcta | 360 |
| tctatntaat | gatgttt | | | | | 377 |
| <210> <211> <212> <213> | 19194 368 DNA Glycine max | c all n locat: | ions | · | | |
| <400> | 19194 | | | | | |
| atgtactgcc | ttangcgatg | ggatgcccat | actaaagcac | aacacgttct | ttcgagcaag | 60 |
| gagtaattca | tctcacaggt | cgtgaacttc | ttacttaggt | agtaaacagc | gcgctctttc | 120 |
| ttcccggatt | cgtcatgttg | ccccagcata | caccccattg | actcgtccaa | gattgtcatg | 180 |
| tacaaaatga | gaggccttcc | aggtactggg | ggcataagca | cgggaggatt | cattangcac | 240 |
| tctttgatcc | ttccaaaaag | cctcttgcaa | tcctcattcc | accagtcagt | ttggttttta | 300 |
| cgcaagaagt | tatacaacgg | ctcacaaatg | gcggtgagct | gcgatatgaa | tctggccata | 360 |
| taattcaa | | | | • | | 368 |
| <210> <211> <212> <213> | 19195 423 DNA Glycine max | c c | | | | |
| <223> <400> | unsure at a 19195 | all n locati | lons | | | |
| agatactcag | ctctaaattg | aattaaacat | tcagaaactg | ctggtaatcg | anatcatata | 60 |
| tgtgtaatcg | attacacaag | gcagattntg | aattcaaatt | ataatagttg | ttgtaaatca | 120 |
| gttttggcca | ctggtaatcg | attaccagag | agtaaatttc | ttgtaaaaga | ctttctaact | 180 |

taattttctt ggccaaacct tttgctactt caattggaat tcccttccta tttaatatac 240

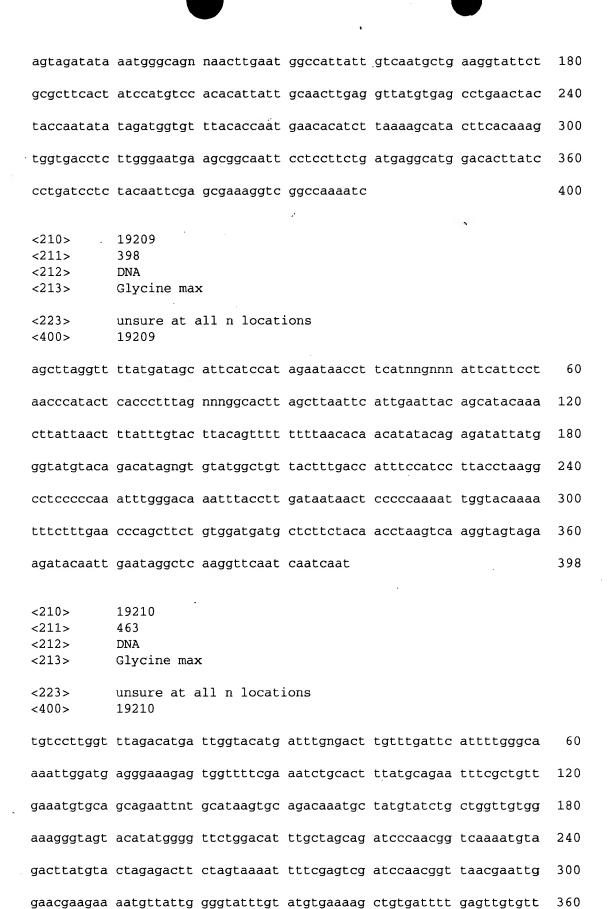
| cctttctaag | actctataga | ctgtcttgat | catccatctt | gaatatcttt | aattgctttg | 300 |
|-------------------------|------------------------------------|------------|------------|------------|------------|-----|
| tcttgaataa | agctgtgaga | cgcatgtgat | ctttnggcat | catcanaaca | tcggcttgat | 360 |
| cctttgtcta | caatctcccc | ctttttgatg | atgacaatcc | ctgaaatcaa | gacaaactat | 420 |
| ata | | | | | | 423 |
| <210> <211> <212> <213> | 19196 373 DNA Glycine ma: | x | | | | |
| <400> | 19196 | | | | | |
| agcttttgcc | tcttcatgtc | tggaatatga | atgttgcata | tagatccaaa | gaaccttagg | 60 |
| tgctttgctg | atggcttatt | cccgttccaa | gcttcaatag | gtgtcttgtc | ttttacagac | 120 |
| ttagttggag | atctgttgag | tatgtaaaca | gcacagtaga | ctgcttcagc | ccacaatgtg | 180 |
| ttatgtactc | tcttctcctt | gagcatcgat | ctaaccatat | ccataattgc | gcaattcttt | 240 |
| ctctctgaca | cttcattctc | gtgaagagaa | tatttgacta | taaggttggc | gctcaatgcc | 300 |
| ttcatcctca | caaaatcttt | catactcgcg | agaggtgtac | tctttgccgg | gatcacttca | 360 |
| ttaaactttt | atc | | | | | 373 |
| <210> <211> <212> <213> | 19197 477 DNA Glycine max | κ | | | <i>1</i> | |
| <223> <400> | unsure at all n locations 19197 | | | | | |
| tcaagaaaaa | gatggcctca | gcaaattcct | tatttccaga | tagtttctct | atcaatagac | 60 |
| ctccaatctt | taatggagag | ggttaccact | actggaaaac | ccgaatgcaa | atttttatcg | 120 |
| aggcaataga | tctaaatatc | tgggaagcca | tagaaatagg | gccttatata | cccaccacag | 180 |
| tagaaagagt | ttcaatagat | ggtagttcat | caagtgaaag | cataaccata | gaaaaaccta | 240 |
| tagatagatg | gtctgaagag | gatagaaaac | gagtacaata | caacctaaaa | gccaaaaaca | 300 |
| taataacatc | tgccctagga | atggatgaat | atttcagggt | ttcaaattgt | aagagtgcta | 360 |
| aggaaatgtg | ggacactctt | cdattaacac | atgaaggaac | tacadatdtt | aaaagatcta | 420 |

| | ngataaatgc | actaactcat | gagtatgata | tatntagaat | gaatgcaaat | gaaaata | 477 |
|---|----------------|----------------------|-------------|------------|------------|------------|-----|
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| | <210> <211> | 19198 414 | | | | | |
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| | cggtgcctat | agtagatgac | gaagcatatc | ttaagacagt | actttatgta | ttaacggtgg | 60 |
| | taaatgaagt | gaatgttatc | ataccaagtg | acaaaattgt | attcgttacc | attggattca | 120 |
| | atgttcaaaa | gtgtcattca | aataaaccaa | agcaaaactg | ccaattcatg | aacaatggga | 180 |
| | tgatggcagc | cgaaatgaag | aacgtatgct | ctgaatggta | ctagaacaca | tgtccttaag | 240 |
| | tatggcagta | tggtgcacgt | ctttctgcaa | gacatttatt | tgagagatgc | ttttgggtaa | 300 |
| 1 | tagggaaatg | atggcagctg | cacggaaata | taaaatatgt | caacatcatt | ctcattcatt | 360 |
| • | gtattgagtt | taggtggtcc | ctaatataat | agggtgcacg | acatttgcta | tcac | 414 |
| | <210> | 19199 | | | | | |
| | <211> | 331 | | | | | |
| | <212> | DNA | _ | | | | |
| | <213> | Glycine max | • | | | | |
| • | <400> | 19199 | | | | | |
| ć | agcttttgat | tattatatcg | acttacatgt | aattcatgag | tcataattta | gtatgttaca | 60 |
| ç | gtacatttta | ggcaatacta | actactataa | aataacatta | atgtgtgcca | ccaagcactt | 120 |
| ć | atagtgaact | tcaatgaaaa | cttttcattt | attaacttga | aaccggcatt | gttaaaacat | 180 |
| t | tattgacaa | gagtgctggt | aaattttcta | tatgatacag | aatcgagagg | attaaactca | 240 |
| ć | agttgcactt | tttctttctt | acttatacta | ttagcatcaa | ttctgttgat | gcaactggct | 300 |
| t | tgaţgtttt | gatgatgaac | aagatgatgt | g | | | 331 |
| < | :211> | 19200 485 | | | | | |
| | | DNA Glycine max | | | | | |
| | | unsure at a 19200 | ll n locati | ons | | | • |
| t | gatattnta | ttgtttcaat | caaagtcata | aagttcatga | tgaattgtat | caaatagaga | 60 |

| aagaagtaa | aa agctggcaaa gaaagcaaca attgatcaaa tggattgaat gaatctcatt | 120 |
|----------------------------------|---|-----|
| gaaagaaat | g aatgatgaat acatgtgttt catattatat acaatcaact ttgggcacca | 180 |
| atctaacta | ac tagaaaatac aactaactat aactgcttct aactgtcaaa acagaaaaca | 240 |
| gttagtgca | t aactaattet atatgttgag ageeeteeea atatgaggaa tgtatgttag | 300 |
| acatttcca | a cttgagctgc aaaagacaaa aagcaatagg agagaaagct ttggtgaata | 360 |
| tgtcggcaa | g ttgatagget aaagatatag atagaagatt gatcaageet gaaagcaate | 420 |
| tcttgcaca | c aatgttggaa gttaattttg atgtgctttg tgcgctcatg aaacacaggg | 480 |
| ttgac | | 485 |
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| <400> | 19201 | |
| gcaggtaaga | a actegegeta eceggettge aaccaacaca etggageaaa ggegatageg | 60 |
| ccaccgacaa | a tgcatgaccc acggcgtcac gcacacggtt aaagagcg | 108 |
| <210> <211> <212> <213> | 19202 213 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19202 | |
| ttttctatag | aattattaaa caagcactta caagtacaac acagaggttt acctgtgatc | 60 |
| gaagaacttc | cttgtatgta ccaatttctc gtaaggctat ggtgaatctg gtcataacag | 120 |
| gacctaggct | ataaacgtcc gtcntanatc atttcatcac cattagactt ttagagtcaa | 180 |
| aggcaacata | cagactctaa atcaaattat gaa | 213 |
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| <400> | 19203 | |

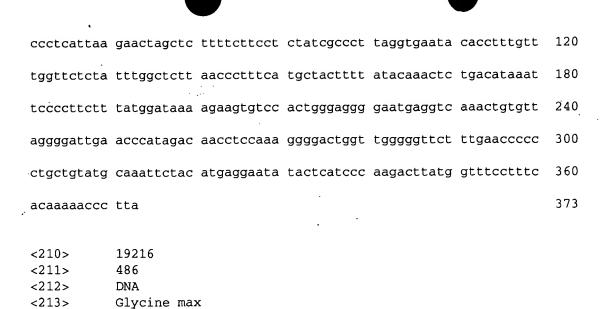
| tatgccca | tg gacatgttga tatgaccaac aagtatgatt atgtcctaaa cctagcgcct | , 60 |
|-------------------------------------|---|-------------|
| atgcgtgc | ga aagggcatcc aatgtcaatg acaagatatc caaacaaaaa tgtggaagct | . 120 |
| actacagt | gt cetttecatg tactaaaaat atcaagaate cagegettaa catageaage | 180 |
| gaaaaacto | ca tttgcgacag ctagccacct tgtcatatga | 220 |
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| agcttgago | t tattgttgct gttccacaaa gctccacgga atttggctcg gccatgctct | |
| | | 60 |
| | g ccctcttggt ttctttttca agggctcttg cggtagcttc attttcttct | 120 |
| | a cacactettt ecagaegtet atagegaeta aettgaattt ttetttggea | 180 |
| agtcttgcc | t ttcctagatc tgtttttaga gcttggactt cttcatcctc ttccggagct | 240 |
| tcgaagttc | t cctcgttgat aattttcaac ttggagagcc aatctaaccc tcgtgtacga | 300 |
| actttcagt | c attcatgata accaccaatg atgccataac ggatgcccct aaattcttta | 360 |
| tettttetta | a acaggc | 376 |
| <210> <211> <212> <213> <223> <400> | 19205 484 DNA Glycine max unsure at all n locations 19205 | |
| cgtagaatgg | | |
| | ctagacatga tacatgtcag ggtttggttt ggttctatgt taatattgat | 60 |
| | atttccatga cacaaatgca aaaatgatga tttggaaatt ntatgcaaaa | 120 |
| | atgcacctat gtggacgctc aagtgtcaat ttttatggtc atgtgatgct | 180 |
| agggctcang | atteatttee tetattntaa ateaaceeaa tgttteeaaa atatgttett | 240 |
| ttatcaactt | gtgcattcat ccaagtccat ttcgggcgtc cggagaaatt atcacagcat | 300 |
| tcacccttta | ggtgtagaca cgtttctttc tttttcataa atcggttatg atcaatgaat | 360 |
| tctttttca | aagaacagtt ggaaatcatc tcttttcata agcatgtcgg tatctagcta | 420 |
| | tttctctttg tcaccttgtt gttachtan | 480 |

| tttt | | | | | | 484 |
|----------------------------------|------------------------------------|-------------|------------|------------|------------|-----|
| <210> <211> <212> <213> | 19206 380 DNA Glycine max | x | | | | |
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| agctagagtt | tttctttata | tatgacatgc | atgatgccct | tťcccactgt | atccacttaa | 60 |
| atttccatat | gctagaaaat | cattaatagt | acaaaacacc | attgtgcgta | acctgaatgt | 120 |
| ctactgcaca | tttgcatçcc | acacatctac | cccttcttcc | cacaattgtt | tcaagtcttc | 180 |
| gattaatggc | gtaagataca | catcaatatc | attccctggc | tgccttggac | ccgcgatcat | 240 |
| catacacagg | ataatgtatt | tacgcaaaat | gcacaaccat | gggggaaggt | tgtaaatcat | 300 |
| cagtaaaaca | ggccaggaac | tgtggttgct | gcttaagcta | ccataaggat | tcattccatc | 360 |
| agaagcaaga | gcaagcctta | | | | | 380 |
| <210> <211> <212> <213> | 19207 225 DNA Glycine max | | ions | | | |
| <400> | 19207 | | | | | |
| tgtgactctt | ggcaatntct | ttaaaactag | tcacttacaa | agttgtgatt | tttgtattaa | 60 |
| tcttcagaaa | caagtcactt | gaagaattgt | gacttttgga | aatttattt | tcaaaatcag | 120 |
| tcactggtaa | tcgattacca | ttaaggtgta | attgattaca | catgaacaga | tgtgactctt | 180 |
| cattttaaat | gttgaaaatt | aaaacgttaa | tatgctctgg | taatc | | 225 |
| <210> <211> <212> <213> | 19208 400 DNA Glycine max | | | | | |
| <223> <400> | unsure at a 19208 | ll n locati | ons | | | |
| agcnagnnat | ttatcttatg | attaatcaag | aacaagcttg | agcgcacatc | gttcgcgtgt | 60 |
| actatatcca | ctcgacaagg | tttgaagtac | aggagacctt | caatcctata | acgcaattct | 120 |



| ttgggcagag | tatctgactt | tgccctgttt | cgcttggttc | tgtagtccat | gatgattgga | 420 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| tgtggaatta | cctggatgtt | gtggatagct | tgggaggatt | gat | | 463 |
| <210> <211> <212> <213> | 19211 381 DNA Glycine max | ĸ | | <i>:</i> | | |
| <223> <400> | unsure at a | áll n locat: | ions | | | |
| agctngccgt | tatggtgtgt | ttcgactatg | ctctcgggtg | gcggaacaag | ctacaaaagg | 60 |
| agagagcaag | aaatgaatag | ccaatggttg | atacatgggc | ggagatgaaa | aggatcatga | 120 |
| ggaagcggca | tgtgccggct | agctactcaa | gggatttgaa | attcaagctc | taataactaa | 180 |
| cccaaggcaa | catgggggtt | gaggagtatt | tcaaggaaat | ggatgtgctc | atgattcaag | 240 |
| cgaagattga | agaagatgag | gaggtaacta | tggctcgatt | tcttaatggg | tcgactaatg | 300 |
| atattcgtga | tatcgttgag | ctgcatgagt | gcgttgägat | ggatgatctg | cttcacaaac | 360 |
| cactccatgt | agagcaacaa | t | | | | 381 |
| <210> <211> <212> <213> | 19212 393 DNA Glycine max | ¢. | | | | |
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| ggtgcatcca | ataccctgat | gaggatgtcc | catatgttct | taaatctgta | ctgattcatt | 60 |
| tgcttccaaa | gtttcatggc | cttgcaggtg | aagacccgca | caaacatttg | aaagaatttc | 120 |
| acattgtctg | ctccaccatg | aaacccccag | atgtccaaga | ggatcacata | tttttgaagg | 180 |
| cttttcctca | ttcattatag | ggagtggcaa | aggactggct | gtattacctt | gctccaaggt | 240 |
| ccatcacgag | ctgngatgac | cttaagatag | tattcttaga | aaaaaatttc | cctgcttcca | 300 |
| ggaccacaac | catcaagaag | gatatctcat | gtattagaca | actcagtgga | gagagcctgt | 360 |
| atgagtactg | agagagatat | aagaaactat | ata ' | | | 393 |
| | ggagagacac | aagaaaooao | 5-5 | | | 333 |

| | | • | | | | | |
|---|-------------------------|------------------------------------|----------|------------|------------|------------|-----|
| | <212> <213> | DNA Glycine max | | | | | |
| | <400> | 19213 | | | | | |
| | agctttatct | ttcaatttcg agcg | ctcgt (| tatattacgg | gactcaatca | gacatccaag | 60 |
| | taaaaagtta | tcatcgtttg aatt | gctca (| gagcttcaac | attcaatttc | gaacgactcg | 120 |
| | atatatgatg | ggactcaatc agac | atccga 🤉 | gtaaaaagtt | attgtccttt | gaaatggctc | 180 |
| | agagattcca | cattcaattt cgag | egtete a | aatatattac | cggactcaat | cagacatccg | 240 |
| | aaaaaaaat | tattttcgtt tgca | ttgct (| caaaggttca | acattcaatt | tcgagcgtct | 300 |
| | tgatatatta | cgggactcta tcag | acttcc q | gagtcaaaag | ttattgtcgt | ttggatatgc | 360 |
| | ttcaa | | | ·** · · . | | | 365 |
| , | <210> <211> <212> <213> | 19214 454 DNA Glycine max | | | | | |
| | <223> <400> | unsure at all n 19214 | locatio | ons | | | |
| | ctgagacaat | tcatacgaca ataa | ctgtnt a | actcggatct | ctaatttagt | tccgtaacat | 60 |
| | atcgagatgc | tcgaaattga atgt | ggaatc t | tctgagccaa | ttcaaacgac | aataagttnt | 120 |
| | tactcggatg | tctgattgag tccc | gtaaca t | tatcgagacg | ctcgaaagtg | aatgttgaag | 180 |
| | ctctcagcca | attcaaacga caata | acttt t | ttactcggat | atctgattga | ttaccgttat | 240 |
| | ataacgagac | gctcgaaatt gaatq | gttcaa (| cctctgagca | aattcaaacg | acaataactt | 300 |
| | ctttctcgga | tgtttgattg agtco | ctgtaa t | tatatcgaga | cgctcgaaat | taatgtttaa | 360 |
| | gctctatcca | attcaacgac ataad | ettta o | ctctatgtct | gttgagtcca | taatatacga | 420 |
| | gacctcgaca | tgaatctgaa ctcta | attcat t | tcaa | | | 454 |
| | <210> <211> <212> <213> | 19215 373 DNA Glycine max | | | | | |
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| | agcttgtatt | ttcttttatt atgga | attga t | tccttcctaa | gatggagcca | aacccactca | 60 |

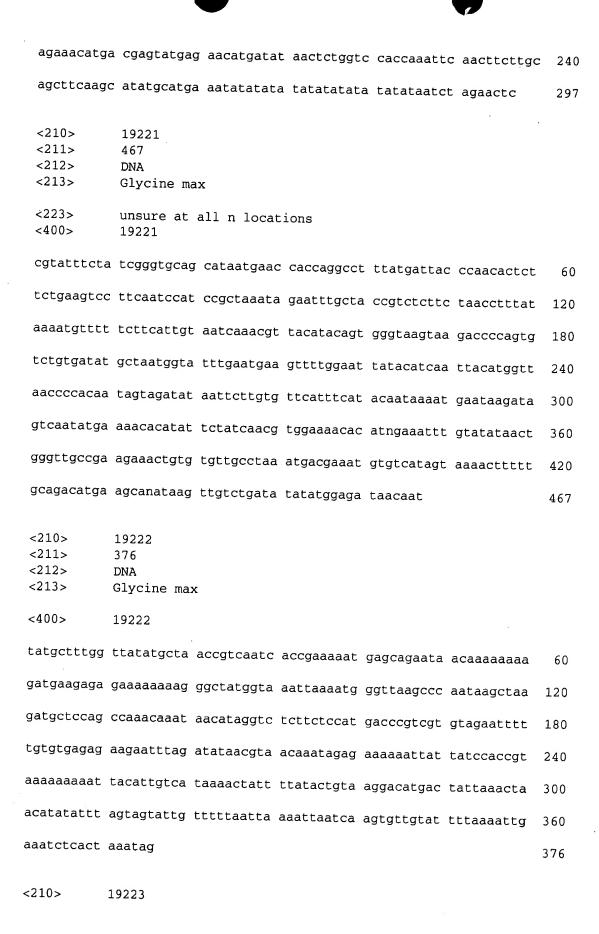


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|----------------|----------------------|--------------|------------|------------|------------|-----|
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| gttttcacta | gttttcatta | tatttcatct | acgttggatc | aattcacacc | atggaatgta | 120 |
| atagcattat | attccatttc | cctccattcg | atccttctcc | accaatcgaa | acattagtgt | 180 |
| gtcaagattg | tatgcaataa | agtggcttaa | gtttgagtcc | ataaattgct | ttattcagat | 240 |
| tatatacaag | ggatttatca | gaggactcca | cacaaacatg | gggcagccaa | taacctcctt | 300 |
| attccagacc | tctgaaaact | acagcctctg | acagttagga | tgggaagtcc | atatctttag | 360 |
| aaacttataa | agagaaccaa | aaaccttctc | agtgttatga | aactggcaca | tgtgacatcc | 420 |
| tgagaattct | actcggaatt | tctgtaagta | ttacatttaa | ataattatat | agatatatat | 480 |
| tattcc | | | | | | 486 |

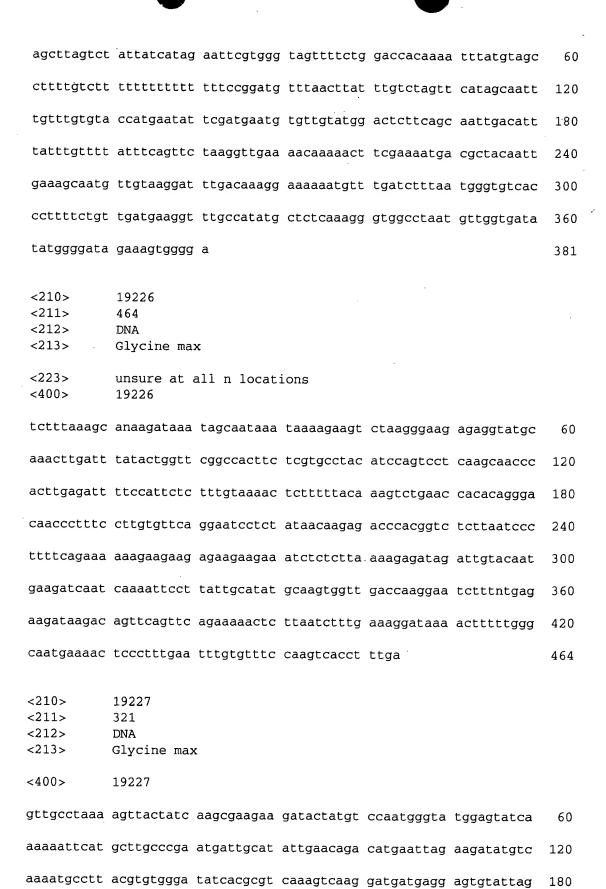
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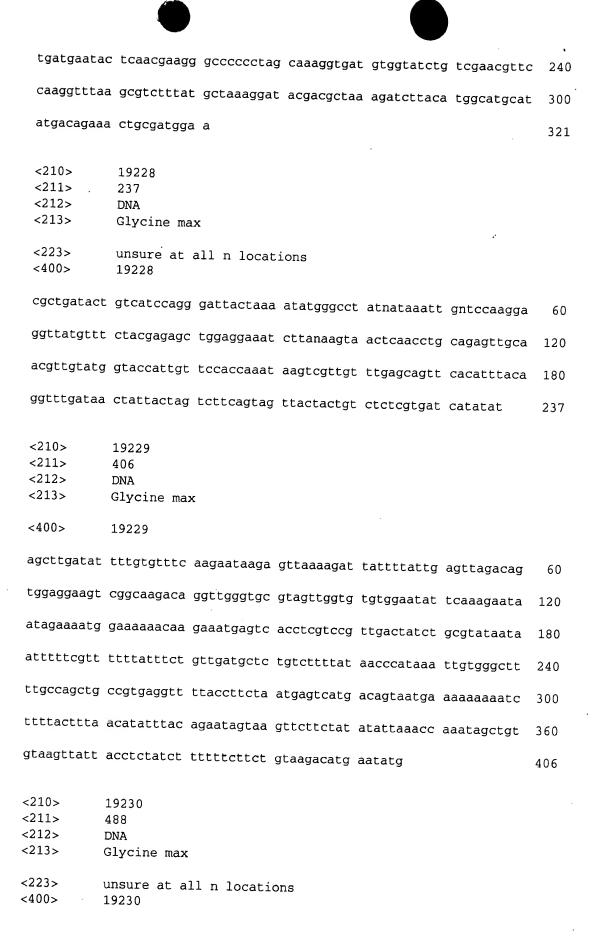
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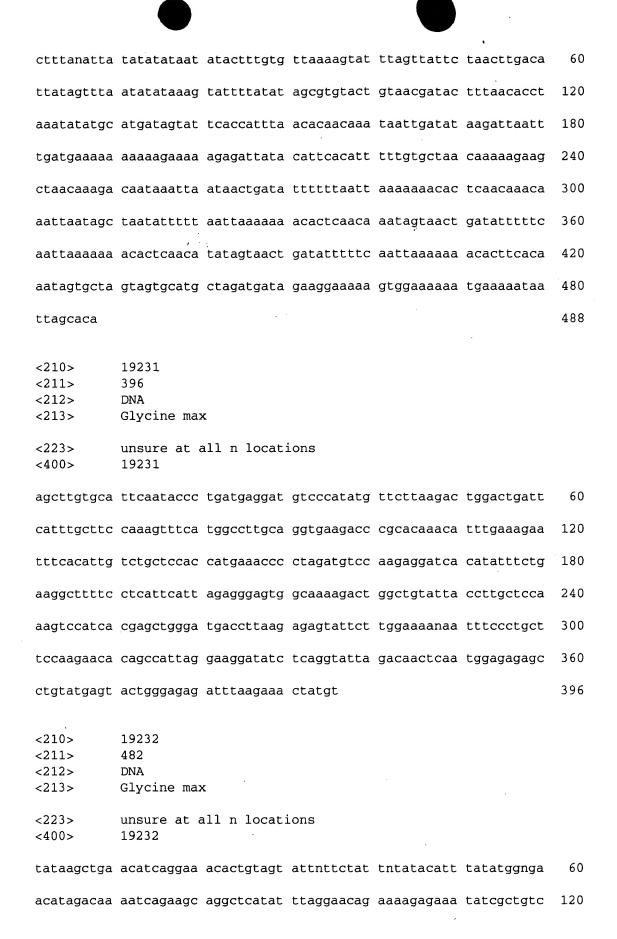
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|----------------------------------|---|-----|
| taaaatcac | cc acttaagatc tatgatgcag tgctcccacc aacggccttg gacatatgac | 360 |
| tcc | | 363 |
| <210> <211> <212> <213> | 19218 358 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19218 | |
| agcttatta | t tattnnnnaa ancnatatga agcaaacaaa taaatatttc actgtaatat | 60 |
| aaactttgc | t tccaattata tatactcgct gaacttctta cattataaat tagaatagat | 120 |
| | t taacaacaac aatatgttgt tattgtcgtt gaccaccata ataagatgat | 180 |
| | a ctctgatctg gaatgaaatt aaacatttca tactaaactt gctcaacctt | 240 |
| | a atgagataat cataatcata ctggacttgg tagacacttt atcatcatta | 300 |
| gtgccgatga | a tggaattgat ttcaaagctg taataagtgg agatctttta ccaccata | 358 |
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| tactaagctc | ttgcgtaccg tgctgggctc acagaatgcc aagtcatttc ttttttacta | 60 |
| gctatgtgaa | ctgaatgatc ctgacgtacg acctt | 95 |
| <210> <211> <212> <213> | 19220 297 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19220 | |
| agcttttgct | tatggatcat tcattggngc acgtgacagt tatatcatac agctatactc | 60 |
| atatattgtg | tgatcatgta cttgtttaca tcagctgaat tcttgtgatt ctattgatat | 120 |
| ctacttgatc | atttgccagc attgtagggg ctctatttct aaggcctatg caactggtga | 180 |

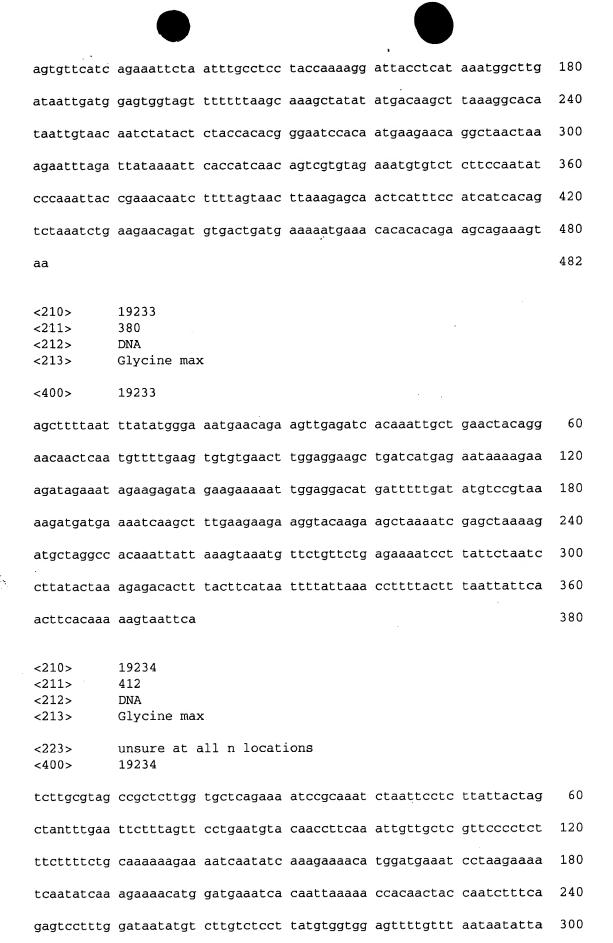


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|-------------------------|--|-----|
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| attgctagtt | agcatgcaat taatctcagt tgtgttttca caggataaaa ttcaaagaca | 120 |
| ttgggagcat | cttgagcttt tgttttcaat agtcatgtca ataatatttg aaagggggtc | 180 |
| tttggaaacc | gagaaaaatg tcaatatgct ttcttcaaga attcacaatc ctttatgtaa | 240 |
| attaagtggc | tttcctaaag gtattgtgga gaagaatcaa atcaacaatg ttgacttatc | 300 |
| accaacaaan | agcattanga ttccacacac ggataaatta ccacaatata catcttgggt | 360 |
| atacgtggct | aggtgcgtct gactntcaag tttcattttt tttttgtcca atagatgacc | 420 |
| tacgaattgc | aatgcaattt tgtcataaga taaaatacat gcttcttga | 469 |
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| <400> | 19224 | |
| tcttcttttg | ctgttgctat ctgattgtcc atgtcagcca gtttttcagt tggttcaatg | 60 |
| tttcctgaat | caatcaaggc cataatette teeegggeag catetggate taettetata | 120 |
| tgagcacaga | caaatatete ttegagetet gettgettet tgaaageaat tteetteate | 180 |
| ctgctggctt | tcagctgatc aagtctctca acttccactt cagcctttat tattaaagga | 240 |
| aaaaaacagt | taatatcata cttggaactc gaatctgtaa acttcataag ttctttagta | 300 |
| cctacctgct | caatcagatc cagagcaagg gcaccaggaa cagtgacttc atcaacagaa | 360 |
| gctgacatat | tacaggtaac atggtcaaat agtctccttt cctcgggatg agtatccatt | 420 |
| agattccaaa | gatcaattaa ctgagaagct aattcttg | 458 |
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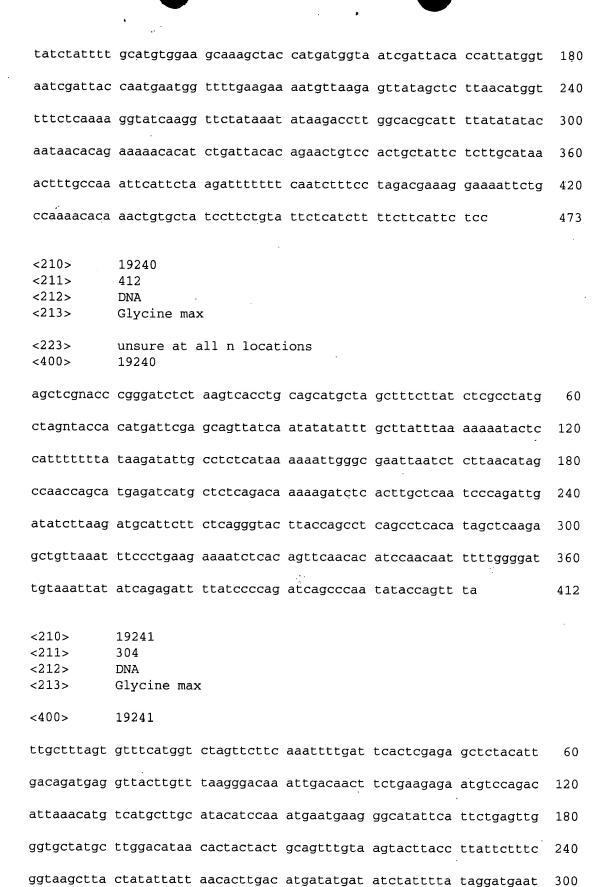




| tacttttgcc | ttccaaaaaa | aacttatgac | tgatcctctc | ttcattaatc | ctattttaga | 360 |
|----------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| tgttattgta | taaaagatca | taggttctcc | acctgcctac | actattcctc | ct | 412 |
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| <400> | 19235 | ÷ | | | | |
| tttattttat | attcctcact | agcttatttc | tcaatgtatt | gcaccgaact | taaatttata | 60 |
| ttttattatt | tataggaata | tttttaatat | attattaatg | atttagtgta | aattttgaag | 120 |
| gatatgccaa | taaatcattt | atcaaattct | taggtattat | aatcgattta | ggattctatt | 180 |
| aaaaataaaa | taaaatttaa | taaatataaa | ttttatatta | attaaattat | ttagaaaaaa | 240 |
| taaatataaa | gtatcatgtc | acatcactct | cataaaataa | aatcttatat | atatatatat | 300 |
| atatatatat | atatatatat | atatatatat | atatatatat | atatgcacaa | gtttagtgtc | 360 |
| tcataccttg | cata | | | | | 374 |
| <210> <211> <212> <213> | 19236 350 DNA Glycine max | s | | | | |
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| tagcaacata | cacaaacata | catacacata | ccaagagaca | ctatcctata | ctctatccgg | 60 |
| tcaattgtaa | aacatcatta | atattcttat | caatattacc | atcatcgtca | taagagaagc | 120 |
| atatgaatct | tattcgtcaa | tctgattcac | actacatgat | tcaacacaca | aatgcaatct | 180 |
| atccaaaaat | ccttcaagtc | ttatggttct | tacaacaaaa | ttctgaaaag | ttttctagca | 240 |
| ttgaaaaccc | cccatttttt | ttatcacaag | catctcccaa | tcctccttga | ccaaaaaacc | 300 |
| cacctccaca | aatgtgatgg | ttgacaacta | tgttgttggg | acccccccc | | 350 |
| | 19237 579 DNA Glycine max | . | | | | |
| | | ill n locati | | | | |

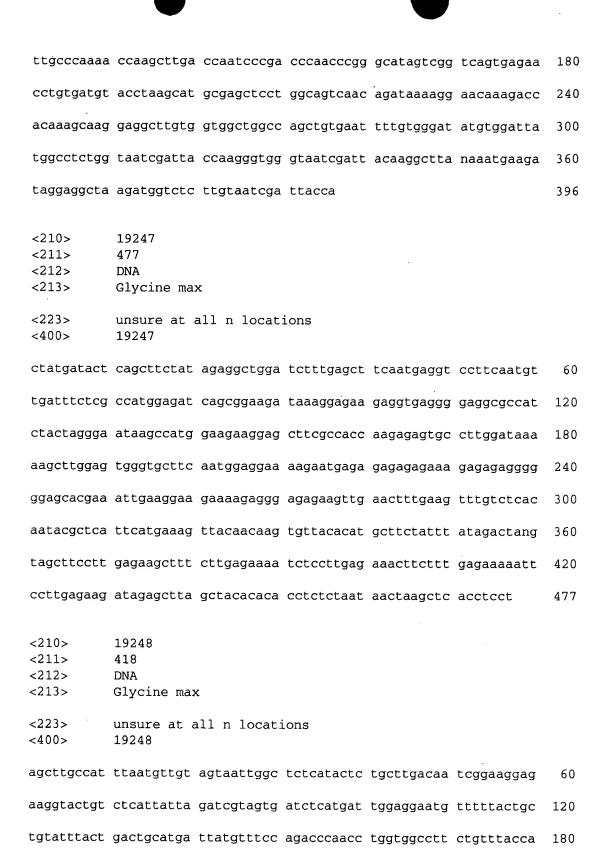
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|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|--|
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| aannnnnaaa | gagagtgcnt | ttgatgcaat | cgataggaca | nacgcgaaca | tananaccca | 120 | |
| agccacagga | gtaacaagaa | cgcttcagga | gaggcgatga | tttcatctac | acgactcata | 180 | |
| agctcaaacg | cgtagctaag | aagatgacga | ccagagcgac | gccaagatag | aggcgagatc | 240 | |
| accgcaagat | gaaattcgcc | ctccagaacc | acgtgtcaag | aggccagaac | cacgacaaga | 300 | |
| cccacgcgag | agaggacaga | tagcaccccg | gccaaaactg | ggagcacagg | agctgcccaa | 360 | |
| accttgacca | aagagcgcta | ctctctggaa | ccgacaccac | aaacgagaag | gacaccacta | 420 | |
| gcaaaaggga | acagaaagca | ttaacagagg | gacgacgggc | caaccgaacg | caacacgggg | 480 | |
| aagaccagac | aaagaagccg | gaaacaacac | caaccggttg | gaccgcggca | accccagcga | 540 | |
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| | | | | ttagttgtac | | 180 | |
| | | | | atacataatc | • | 240 | |
| taccacagaa | tgggtatgtg | taactcatcc | aaacatggct | atttcaacag | gctctcaaca | 300 | |
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| agccgaacat | cttcttaagt a | ataaccagt | gatggcattt | tcatgatcgg | ataagcatct | 180 |
| ctatgacaaa | gaatttgatt a | ataatatca | atattccaac | atctatttgt | agcatcaata | 240 |
| aaattttcca | cctttttatt c | tccacatca | ttaatgagag | gagtttcaat | gaaaagatta | 300 |
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| acaccaatta | ttgttggata g | gtatcatca | tatttatgct | aagaaaatca | tgcacatcag | 120 |
| aaagaataag | ttaaaatagt a | taccaattg | gtacaatagc | accatctgag | ccaccacaaa | 180 |
| gcatcaagtc | ctgcaattac t | tacattntt | aatgttttca | ataaaataac | tatataatta | 240 |
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| agcaaatgct | gaagtgagat a | cttacagct | tcacctctaa | tgatatggtt | tgcagcattc | 360 |
| aatatacaaa | aattactagt a | gcacacgct | gtagagattg | aataaatang | gcccatccac | 420 |
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| cttttgtctt | tttttttt | tttccggatg | tttaacttat | ttgtctagtt | catagcaatt | 120 |
| tgtttgtgta | ccatgaatat | tcgatgaatg | tgttgtatgg | actcttcagc | aattgacatt | 180 |
| tatttgtttt | atttcagttc | taaggttgaa | aacaaaaact | tcgaaaatga | cgctacaatt | 240 |
| gaaagcaatg | ttgtaaggat | ttgacaaagg | aaaaaatgtt | tgatctttaa | tgggtgtcac | 300 |
| ccttttctgt | tgatgaaggt | ttgccatatg | ctctcaaagg | gtggcctaat | gttggtgata | 360 |
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| acttgagatt | ttccattctc | tttgtaaaac | tctttttaca | aagtctgaac | cacacaggga | 180 |
| caaccctttc | cttgtgttca | ggaatcctct | ataacaagag | acccacggtc | tcttaatccc | 240 |
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| gaagatcaat | caaaattcct | tattgcatat | gcaagtggtt | gaccaaggaa | tctntttgag | 360 |
| aagataagac | agttcagttc | agaaaaactc | ttaatctttt | agaaggatan | aactgtttgg | 420 |
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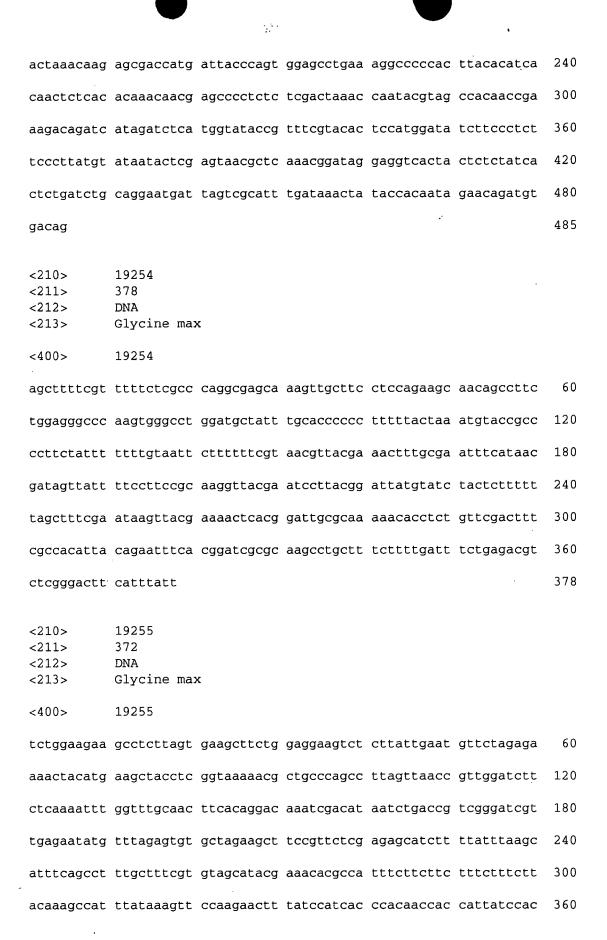


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| tgtctttt | c tgtaccttca cactattcat agcttttaga tctaaatcat tgttaatatt | 360 |
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| ttaattatg | g gcttaacatt ctacagggca ttcaaaatga gaacaaaaac ttaaaaaaat | 180 |
| cactgaaggt | aatttaatat tagatgttac aattataaac ctttctccat catattttgt | 240 |
| ggagatattt | tgcattacta ctggtatcca tctagtttct ttaccttgtc atgatttgaa | 300 |
| ttgtggaatt | gtcacctgta ttctttgaat tgttnttcca ggatgtggtt actgagaatg | 360 |
| aatttgagaa | a aaaacttett getgatgtta tteegeeaae egatattggg teacatttga | 420 |
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| tatttatcaa | taaatatcta attagcaatg attaaatatt ccagattcta tcaaagcatt | 180 |
| tatgagttat | tacgacacag aagcatcaga tgggacttca acattntaac cacaagcagg | 240 |
| tcaataaaac | ccatttaatc tgcaagtcag tgttggatgc ctattcataa tgtgctatac | 300 |
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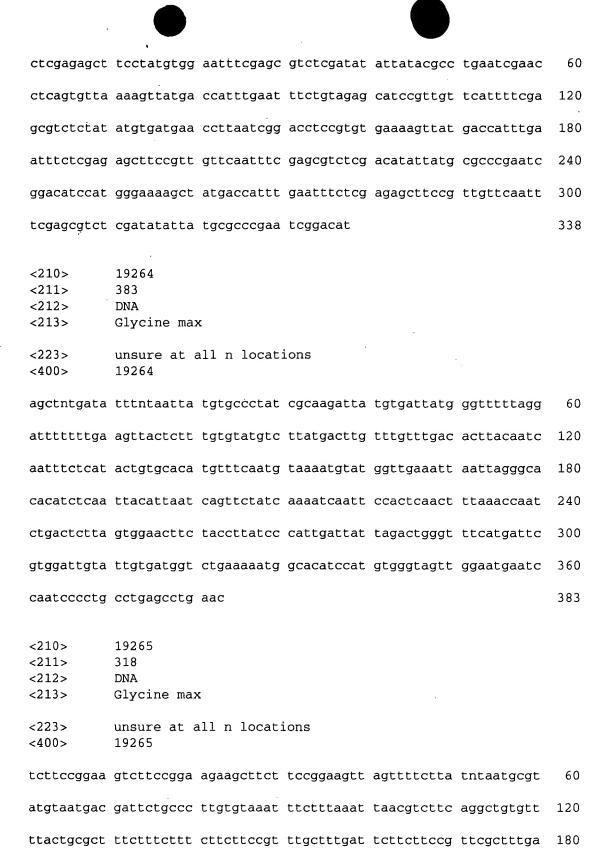
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| cageetetge a | ctagacctt gcaacaacat tttgcttctt actcttccat gagacaagat | 360 |
| ttcctccaat a | gacacacaa tatcctaaag tggaacgcct atcaatgcgt gatcctgccc | 420 |
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| | agtatgccg gaaaaagtgg gggaaaaggc tgacggcgga gtcttccgat | 120 |
| | agatgaacc tettecagaa agttteegga agaagtgtte tteeggaagt | 180 |
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| (213) G1) | ycine max | |
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| atatatatat ata | atatatat atatatata atatatatat atatataaag cttagggaac | 180 |



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| gaacaaatga | aacattgcgc | tcgaaaaccg | aaccgcgcca | gaacttgaag | agagcagacc | 360 |
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| cctacgcgcc | gggaaaaaac | gccacccgaa | gcggggggca | gagaacaata | agaccaccaa | 540 |
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| tctcttgatc | gcttagtgcc | aggaggtatt | tgtatttggc | ggaaaagagt | gacttataac | 240 |
| ggacctggaa | tggccttcca | tgccatggta | agagtgccag | acatgtatca | caacatgggt | 300 |
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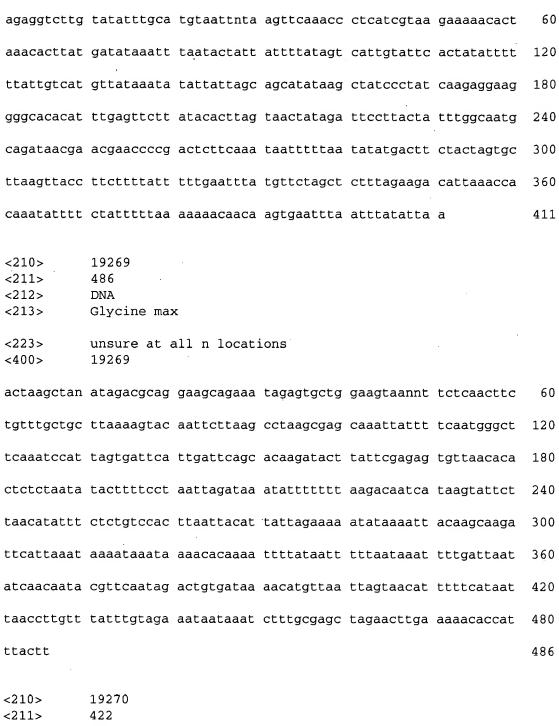


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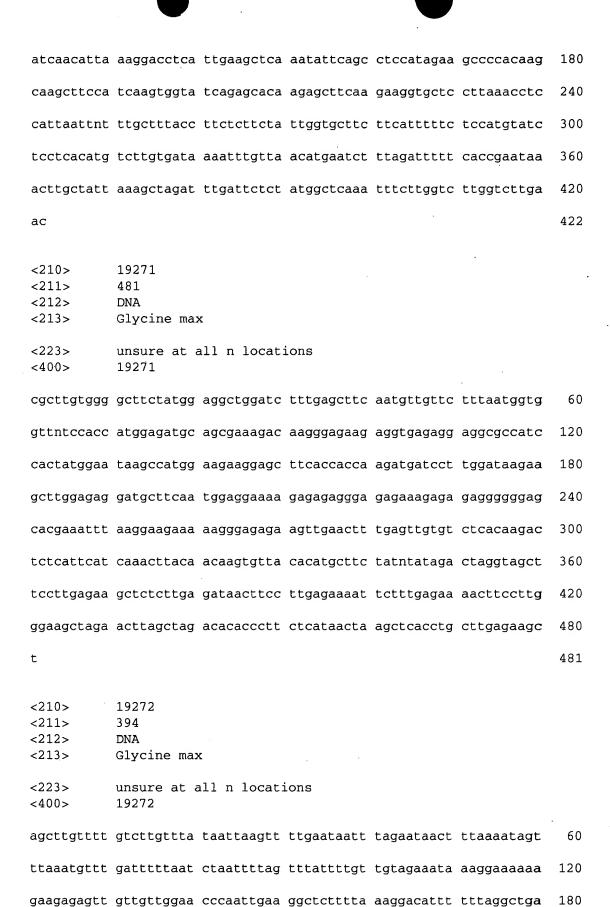
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| tgtacccagt | tgagcgatac | atgaagatct | tactagggta | tacaaagaat | ctatatcgtc | 360 |
| cagaagcatc | tattgttgag | aggtacattg | cagaagaagc | cattgaattt | tgttcataat | 420 |
| acttacagaa | tgctatacct | gttgggcttc | ctgagtgtct | gcatgatgat | a | 471 |
| <210> <211> <212> <213> | 19274 378 DNA Glycine max | ĸ · | | | | |
| <400> | 19274 | | | 2. | | |
| agcttttaat | gtcctatagg | gacaagtcac | aggtggaatg | catggcttta | aaggataaaa | 60 |
| tgaacgcttg | tcaaacgtca | aaaagaagtt | tgaccgaaca | gctgagtaga | acaaaagaaa | 120 |
| atatgttcac | aatcattgac | cagtataagg | aaaatgaaac | ctagctgcta | ttcatgggca | 180 |
| aagactaaag | gatgagcatg | cgaaagtatc | ggctctacaa | atggaaaggg | aagcaagaga | 240 |
| gagagtgata | gaattattgc | acggggaggc | gatgaaatgg | atggatagat | tcgctctcac | 300 |
| tctgaatggg | agtcaagagc | ttccaaggct | gttagccaaa | gcctatgcaa | tggccgatgt | 360 |
| atacccagct | cccgatga | | | | | 378 |

| <210><211><211><212><213> | 19275 97 DNA Glycine mas | × | | | ; | |
|---------------------------|------------------------------------|--------------|------------|------------|------------|-----|
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| tcatcctgag | atccctcttg | ttggactaag | cccaatagtt | attttcctct | taggttcaga | 60 |
| caaacttaga | ctgagtttcg | ttcgcagatc | cttcttg | | | 97 |
| <210> <211> <212> <213> | 19276 414 DNA Glycine max | × | | | | |
| <223> <400> | unsure at a | all n locat: | ions | | | |
| agctttatcg | ttattcccaa | agcttcatgt | agacttgtcc | agaatcgcga | agtgaacctc | 60 |
| ggatccctgt | ctgatacaat | actggaagga | attccatgca | accttaccac | ttctttgata | 120 |
| tacaactcca | ctagcttttc | cattntatac | ttcatattca | ccggaataca | ctgagcagat | 180 |
| ctggtaagtc | gatctacata | ttcagccaca | tctttcttcc | tgccatgcca | ccaaacactt | 240 |
| ctcttcaaat | cttggcacat | cttagacatt | ccacgatgga | aactaagacg | acttttatgc | 300 |
| gcttcttcca | tgatctttac | tgtcaaatca | tctaaagatg | acacgcatat | gctccccttg | 360 |
| aatttaatta | taccagttga | gccctctcga | actctacctc | cttatccccc | atta | 414 |
| <210> <211> <212> <213> | 19277 601 DNA Glycine max | ς. | | | | |
| <223> <400> | unsure at a | all n locat: | ions | | | |
| tacgaacgcg | tacactgcag | naggaagtat | agtcacacct | gatagcanga | gantaaacnn | 60 |
| acaaaacccc | nnaaagagtg | gggtttgatg | ccatcntagg | acacacgnga | caccananaa | 120 |
| cacncaagct | cggagcgnaa | gcaaaccaga | agaacncgaa | gctcggatat | tcgatcgagt | 180 |
| cccgtagtat | atcacgacgc | gcgagcagga | agacagaagc | gcttagcgga | ttaaaacgac | 240 |
| aatgacgatt | aactcggatg | tccgattaag | tcccgcaata | tattgagaca | cacgacagag | 300 |

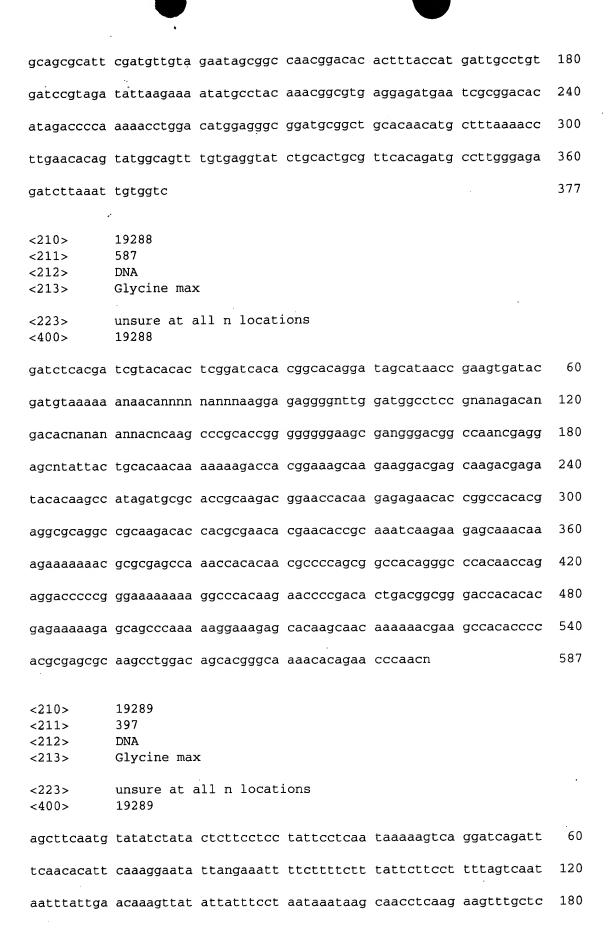
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| gggccgcaat | aaaccgagag | gcgcagaagt | gaaaaaggaa | gcttgcagca | aaggcaatac | 420 |
| acaagaacga | ttaacgcgaa | gatccgatgg | aggaccgcaa | gataccacga | cgctcgaaat | 480 |
| tgacacagag | acactcagca | caagagacga | cactaagaac | tacacgaggg | acgaacgggc | 540 |
| aggcgaaaga | agaatccgcc | gaacggacta | cagaacagaa | agcaacacca | acaggccgac | 600 |
| g | | | | | , | 601 |
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| agctgttacc | tttatcttat | ctcacagacg | ctatatctgg | gagccgatac | agtccttgtg | 60 |
| ttcggactct | cagccactta | cgatagccgc | cgatgatccc | attactgctt | cccgtaagct | 120 |
| ctctgtcctt | tcttcacgcc | gcatcccatg | ccttgcgatc | tccttggagt | actctcgcgt | 180 |
| agtggtcact | agaaccccat | gcgatcaaag | gcctgatgct | tggtgctaat | ggcgctcctc | 240 |
| tcatgggg | | | | | | 248 |
| <210> <211> <212> <213> | 19279 393 DNA Glycine max | ς | | | | |
| | | | | | | 60 |
| | | | | tctctcaatt | | 60 |
| | | | | gcctttgctt | | 120 |
| | | | _ | tctaagactt | | 180 |
| | | | _ | caaggtggac | _ | 240 |
| ttattttagt | gaaaaactat | atagtgccac | cctcaactac | cccacctatg | ataaagagct | 300 |
| ttatgcctta | atatgagctc | tccaaacttg | tgaacattac | cttgttgaca | aggaatgtgt | 360 |

cattcatagt gatcatcagt cacttagcac att

| | | | | • | | |
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| <213> | Glycine max | c | | | | |
| <400> | 19280 | | | | | |
| gacattatca | ttattacttc | ttccacggtg | ctggaacgta | cttacatgga | cttgatgggg | 60 |
| cctatgcaac | tagaaagcct | tggaggaaaa | aagtatgcct | atgtggttgt | ggatgatatc | 120 |
| tctagattta | cctgcgtcaa | ctttatcaga | tagaactcac | ácacctttga | agtattcagg | 180 |
| atgtgagtct | tacacttcaa | agagaatagg | accgtgtcat | caa | | 223 |
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| <211> | 438 | | | | | |
| <212> <213> | DNA Glycine max | ζ | | | | |
| <223> | unsure at a | all n locati | ions | | | |
| <400> | 19281 | | | | | |
| cgtccttggt | ttagacatga | ttgatacatg. | atttgcgact | tgtatgattg | aatttgggca | 60 |
| aaattggatg | agggaaagag | tgattttcga | aatctgcact | ttatgcagaa | ttttgctgtt | 120 |
| gaaatgtgca | gcagaatttt | gtataagtgc | agaaaaatgc | ttgtgtatgg | atggttgtga | 180 |
| aaagggtagt | acatatggag | ttctggacat | ttgctatcag | atcccaacgg | tcaaaatgta | 240 |
| gacttatgta | ctagagactt | ccagtaaaag | tttcgagtcg | atccaacgat | taatgaacta | 300 |
| taacgaagga | tatgttactg | gcgtatttgt | atgtgaatag | ctgtgattnt | gagttgtgtt | 360 |
| ttgggcagag | atttctgcct | ttgcttctgt | tagcttgagt | ttgttagtcc | atgatgattg | 420 |
| gatgtggaat | acctgaat | | | | | 438 |
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| <211> | 276 | | | | | |
| <212> <213> | DNA Glycine max | ζ. | | | | |
| <400> | 19282 | | | | | |
| tgcaagtttg | cttttatttt | tacaagatgc | acatcggtgg | gcagctacct | catgcactct | 60 |
| tctaacgacc | atggcatact | tgctggcact | aaattgctgg | gagtctgagg | ccatcttctc | 120 |
| aattaaatga | atggcttcag | tatgagatat | gtgttcaaag | gctacaccac | tggcagcatc | 180 |

| tatcatact | atctacatat taccgagtcc t | tcataaaaa tata | aggagaa acaacctgat | 240 | | | |
|--|--|----------------|--|--------------------------|--|--|--|
| | g attgtggggg.caaccgacac c | - | | 276 | | | |
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| <211> <212> | 307 DNA | | | | | | |
| <213> | Glycine max | | | | | | |
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| tttcttcact | ttaggagacg gccattccgg t | gttggagaa gato | caacgac aatgcctaca | 60 | | | |
| agattgactt | gcctagtgag tataatgtaa g | tgccacttt caat | gtgtct gatctatctc | 120 | | | |
| tttttgatgo | agatggcgga gccttggatt t | gaggacaaa teet | tttcaa gaatgaggga | 180 | | | |
| gggatgagga | cataaccaat gaccatgaag ca | actggaatg teec | atgacc atatgcagac | 240 | | | |
| ttatacaago | ccaacgcgtc atagagacac gg | gctggtcat ttgt | atcgct gccattgatg | 300 | | | |
| atgattg | | | | 307 | | | |
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| <210> <211> | 19284 384 | | | | | | |
| <211> | DNA | | | | | | |
| <213> | Glycine max | | | | | | |
| <223> | unsure at all n locations | | | | | | |
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| gttgcgtnta | aatccaagcc gtattggagt ca | atattgaag ctgt | ttctag ccttaggaag | 60 | | | |
| ccgtatctaa | gccgtatctt gtattttaaa aa | ataaaaat aaaa | taaatc ttttttggat | 120 | | | |
| actcctagga | | | 33 | 120 | | | |
| | tactatccgg ccgtatccgt gg | gagtategg tgte | | 180 | | | |
| | tactatecgg cegtatecgt gg | | ggatac gggtgcgaca | | | | |
| ccgacacttt | | ttcacagt agtt | ggatac gggtgcgaca | 180 | | | |
| ccgacacttt | tccttttttc acgtatccgg ac | ttcacagt agtt | ggatac gggtgcgaca tatcaa tggattctct | 180 240 | | | |
| ccgacacttt nttttctttt acatcatttt | tccttttttc acgtatccgg ac | ttcacagt agtt | ggatac gggtgcgaca tatcaa tggattctct | 180 240 300 | | | |
| ccgacacttt nttttctttt acatcatttt tatccaaaat | tcctttttc acgtatccgg acaaggaaggtt atttctaatc taaaatttgggt cttttaaata ag | ttcacagt agtt | ggatac gggtgcgaca tatcaa tggattctct | 180 240 300 360 | | | |
| ccgacacttt nttttctttt acatcatttt tatccaaaat <210> | tcctttttc acgtatccgg acaaggaaggtt atttctaatc taaaatttgggt cttttaaata ag | ttcacagt agtt | ggatac gggtgcgaca tatcaa tggattctct | 180 240 300 360 | | | |
| ccgacacttt nttttctttt acatcatttt tatccaaaat | tcctttttc acgtatccgg acaaggaaggtt atttctaatc taaaatttgggt cttttaaata ag | ttcacagt agtt | ggatac gggtgcgaca tatcaa tggattctct | 180 240 300 360 | | | |
| ccgacacttt ntttctttt acatcatttt tatccaaaat <210> <211> | tcctttttc acgtatccgg ac aaggaaggtt atttctaatc ta aaatttgggt cttttaaata ag tggtgggccg tgac 19285 389 | ttcacagt agtt | ggatac gggtgcgaca tatcaa tggattctct | 180 240 300 360 | | | |

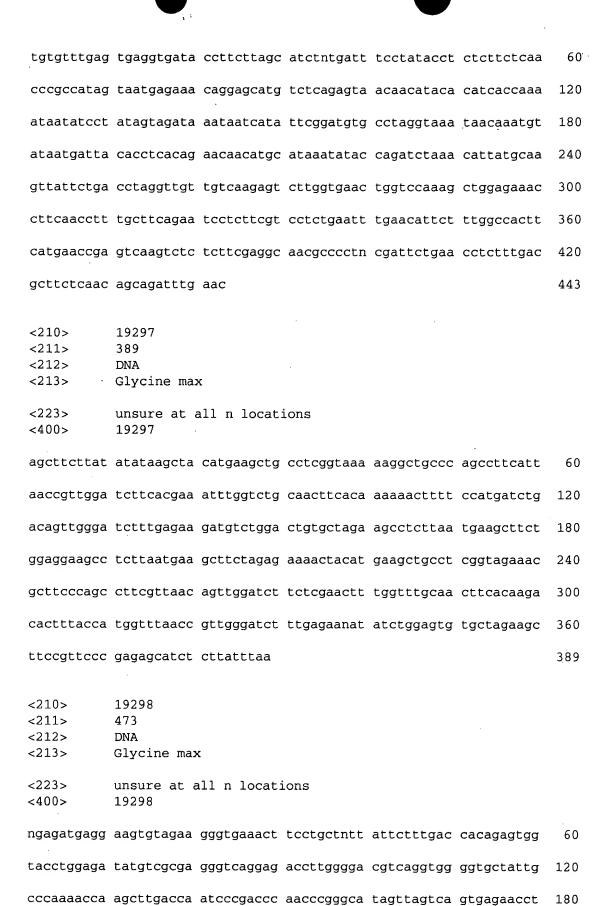
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| tagattetea | tttgaaaaca | tgctgtctgt | ttgacttana | taacaggatg | tgattggatc | 120 |
| atttgtgcaa | gtaacttgca | ı ctttttatga | ttgtcttttg | caccataagg | tgttatggac | 180 |
| aaactcttaa | agaagagato | : tatctcattt | gttatgttct | tttgcaatca | agaaatacaa | 240 |
| ttctttcttt | tctctatttt | ctaatccagc | ttcttaacaa | aatgtgttgt | aaaatttgtt | 300 |
| cattgagaaa | gtagtgtcaa | atctaagata | acatgtttct | attggcccat | gattacactg | 360 |
| ataatatgca | nggatagtat | acctaattc | | | | 389 |
| <210> <211> <212> <213> <223> <400> | 19286 489 DNA Glycine ma unsure at 19286 | x all n locat | ions | • | | |
| acctatgata | ctcagctgtt | ctatagacta | aatatgaaac | tcacctatgt | tagactttat | 60 |
| taatcattaa | gtgagtcttg | caagttacca | aaaaataaat | gatattacat | tnttcaaatt | 120 |
| tctgataact | cgttggattt | aattttcacg | tgatttttc | atgtttatgg | ttctttttta | 180 |
| tttgaacgtc | ccttagcttc | tacggtttca | tataactaaa | tcaaacaaaa | caaatttcac | 240 |
| atctcgaatt | gataaaaaag | gattcaaaat | tgaaaatgat | cattgtttca | gaagaattat | 300 |
| tttggctcac | ttaacaaata | tgaatgatta | aattcaaatg | taaaaaaata | agatatcata | 360 |
| ctaaatataa | tcaaatatat | aaaggactaa | aagtatatct | taacctatct | tttataataa | 420 |
| ataattattt | ttattatata | nattatattt | ttgaatttaa | aaacataatg | aatcttatta | 480 |
| tctaaatat | | | | | | 489 |
| <210> <211> <212> <213> <400> | 19287 377 DNA Glycine max 19287 | ĸ | | | | |
| aaccggagag | aacaacgcat | ggaaggaaga | attgatgtta | gatgacacat | aaccaaagca | 60 |
| ttagaccaaa | agatgacgat | acagagaatg | gattttgaat | tgccgaccac | gcctacagtt | 120 |

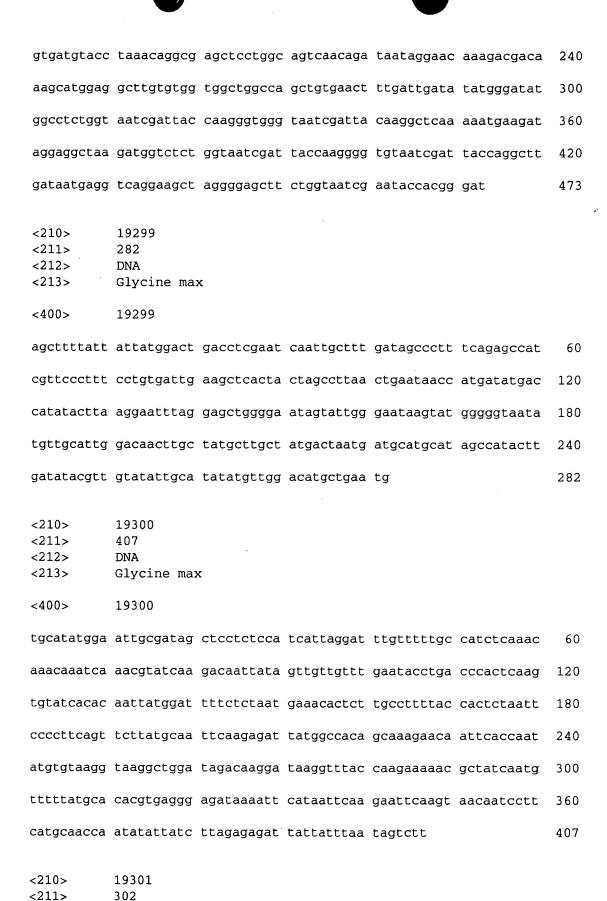


| aaacacatac | caaggagtca | aggactctaa | ttnttagagg | aatcgaagga | tatatatgaa | 240 |
|----------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| agatatatat | gggtattcct | gagagactgt | ttaaagttat | ttgcaagaat | aacaaaaagg | 300 |
| cataccaaat | catgcctttt | cagttccctg | gtaatcttct | tagtattgac | ccattcatac | 360 |
| canagatttc | ttttcagtac | aaagttgcaa | caaactg | | | 397 |
| <210> <211> <212> <213> | 19290 459 DNA Glycine max | x | | | | |
| <223> <400> | unsure at a 19290 | all n locat: | ions | | | |
| tgcttctaca | ggatgggcaa | tggttgaaga | aggacgcaca | acctctataa | gatgatcgta | 60 |
| ctcccctcc | tcctcctcaa | tgagatgact | cctcagccct | catgaatgaa | gttctttcaa | 120 |
| agttacgagg | cctccaaacc | tatgttggtg | aacgctntga | ttccttgaat | ggtcgtatcg | 180 |
| atgccattga | tgctcgcttt | gaaggaatgg | atacccgcat | cactcagctt | gaggaggatg | 240 |
| tgagttatct | tcgtcggtgc | ttcgactttc | ctccaccatc | ttcatagata | tagagtatta | 300 |
| ttatctttga | ttntacgcca | tgtaacgcca | tgtatttggc | tatgtgtttt | aagtcattat | 360 |
| tctttgaact | tagttatatt | tcagtatttg | gcacttatgt | atttatgact | ngaatatatt | 420 |
| taattttgac | ttatgaatgc | gtatgtgaac | tttattata | | | 459 |
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| tgatcttctg | gactaatgtc | attttgtatg | tattctaaga | ttgacgtcat | ctagttttct | 120 |
| tgcctttcta | tcatcatgaa | gaatcttggg | gtatgtatca | tttgacatat | aacactttnt | 180 |
| atttctttgt | gctcttcaat | attagatacc | tttgacaaaa | gtcatctctt | gcattttctt | 240 |
| ctcttggaat | gtaggtgaac | tccatatttt | ttaaattttt | ctttgattac | cccaattttt | 300 |
| tgatgtattt | ttgcaagatt | ggttctttga | cttggtattc | tccattcaat | tggttgagcc | 360 |

| aacaagttaa | gttcgttgga | cactnttagc | actttgacct | ctatgtaagt | actatttct | 419 |
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| <210> <211> <212> <213> | 19292 434 DNA Glycine mas | x | | · | + 8 ₁ , | |
| <223> <400> | unsure at a | all n locat: | ions | | | |
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| tacttgggaa | tcaatcacaa | ataaaataat | gaagtatagg | aattacaaag | tataaggcat | 120 |
| aaccaataaa | aatcataaat | acgacataac | caaaccagaa | tccaaacagt | tcaaaattca | 180 |
| aaaaccacat | agtatcaaag | cataaaagtc | tgaaatccaa | atactgcaag | ataaataaag | 240 |
| tactgaacat | aataatctaa | gtagcatagc | caaataagag | acatagaatt | agaaactaaa | 300 |
| ttctaagaag | gtagaggtgg | tggtggaaga | tcgaaactct | gacgaatgta | acccacatcc | 360 |
| tcttcaagct | gtgtgaggcg | aatatccatg | ccggcaaaac | atgtatccaa | tgagtcgaaa | 420 |
| cgttcaccaa | cata | | | | | 434 |
| <210> <211> <212> <213> | 19293 436 DNA Glycine max | | | | | |
| <223> <400> | unsure at a | all n locati | ions | | | |
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| aaatcttcat | aaataagtca | cttgaagaat | tgtgactttt | ggaaatatat | ttttcgaaat | 120 |
| tagtcactgg | taatcgatta | ccattaaggt | gtaatcgatt | acacatcaac | agatgcaact | 180 |
| cttcattttg | aattttgaaa | attaaaatgt | ttagaagctc | tggtaatcga | ttacaagtat | 240 |
| tttgtaatcg | attacacaag | tttaaaatac | tttaaaactg | tttaaacata | agttataact | 300 |
| cttgaaattt | gaaatattac | cgttttaaga | cactggtaat | cgattactgc | cttctggtaa | 360 |
| tcgattacca | gagagtataa | ctctntggta | atgattctgt | ganaacttct | tatgctactc | 420 |
| aatattctgg | aaaaac | | | | | 436 |

| <210> <211> <212> <213> | 19294 423 DNA Glycine max | |
|----------------------------------|--|-----|
| <223> <400> | unsure at all n locations 19294 | |
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| tcgcatgtac | acttgtaact ccaaggtatc aaacctttca ccaacaaagg tttgaagacc | 120 |
| atcaaacctg | tccaaaatct tntgaagaag agaggaatct tctccaccat gtaagtgtcc | 180 |
| ttcttcatcg | atgggttgag cacccttttt cacccaagag ccatcatgct ctttacggta | 240 |
| accaaaggat | gcaatcactg cagcacctat tagagaggat ctcttgattg gaacataagg | 300 |
| ttcaaaatca | agagggatgt tgaagtgttg aatatagagg gtgactaggt gtggatatgg | 360 |
| caatggagca | tttaatcgca atgccttatg catgagatat cagactaagt gttcccaatc | 420 |
| aat | | 423 |
| <210> <211> <212> <213> | 19295 379 DNA Glycine max | |
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| agcttttagt | tgtttgaaat gtctggcata gcgtcgaaga agtaattatt aacggtttac | 60 |
| tgaaacttga | atgttgtcgt aactaagttg ttgtgcaaat gttgtcttgc cctaacaaag | 120 |
| taatgtttga | atgtatgact atatataatt ttactttgcg tccacacacc atatatgatt | 180 |
| tttcatgttt | actgagaagt taatagatac atactgcttt ttagtgtcaa taacttagat | 240 |
| aataatgtgt | aatttcacaa aacttcaaga gtgtgaaggg aatttgctct aataagtata | 300 |
| actttggcat | tacaaaggtg attatttgtt tttgaacgag ttctcaaaca gcattggaga | 360 |
| atgtagctat | gattgggat | 379 |
| <210> <211> <212> <213> | 19296 443 DNA Glycine max unsure at all n locations | |
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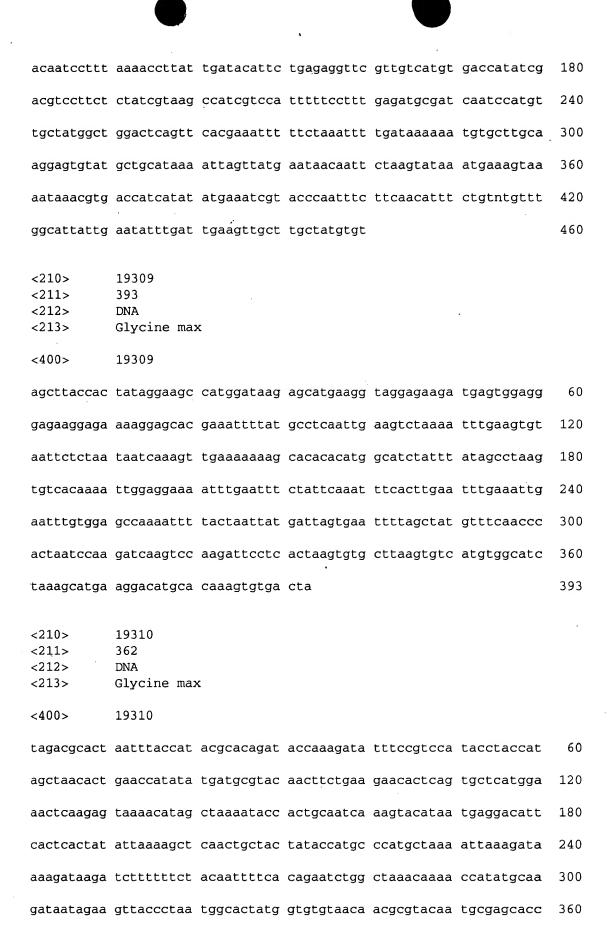




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| ttccacatct | acactttctg aattcatgta | taacattact | aacaaaatat | atatatatat | 120 |
| atatattgat | agtgaatatt tattgtataa | tttaaattta | taaaaatatt | ttatgcatta | 180 |
| gatcatggtt | ttataaaata aattatttgt | tttacaatct | gggatgtgaa | agtagatttc | 240 |
| ctatcatgtc | taatagaatc atgtttttag | agggtcaatg | tagataagca | catatatgga | 300 |
| cg | | | | | 302 |
| <210> <211> <212> <213> | 19302 480 DNA Glycine max | | | | |
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| aacaagtttt | ccacatgcac aatgcgcgca | taaacccacc | atccgctgtt | gcccacctgc | 120 |
| aactgagctc | acgtactccc acgtagccca | tatcctcgtt | gctctcaaca | ccgggtcccc | 180 |
| atcaatcctc | ccaagcttcc acaacatcca | agcaaaacaa | cattcaaaca | gcacaagcta | 240 |
| tcacagccaa | gcgaaacaga gcaaaggcag | aaaactctgc | caaaacacca | accaaatcac | 300 |
| aactgttctc | acttagagac cccagtaaca | attccttcga | tccaattcgt | taaccgttgg | 360 |
| atcgactcca | aaattgtact ggaagtctat | agtacatgaa | cctacattgt | gaccgttggg | 420 |
| atctactagc | atacatacag aactcattct | ggactactct | ttgcacagcc | aaccacacac | 480 |
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| • | atgacgtgaa atatcctgta | | | | 60 |
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| ttacttcgaa | cgatcaattg | agaatatagg | tctgattcgt | acatattgat | , atgagcaggc | 180 |
|--|--|---|--|--|---|---------------------------------|
| gagtatttcc | С | | | | • | 191 |
| | | | | | | |
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| ccccacatta | tttccatgac | acaaatgcaa | aaatgatgat | ttggaaactt | tacgcaaaac | 120 |
| tggtcatgca | tgcacctatg | cggacactca | agtgtcaaat | ttttatggcc | atgtgatgct | 180 |
| agggctcagg | attcgtttcc | tctattctaa | tcaacccaat | gtttccaaaa | tatgttcttt | 240 |
| tatcaattcg | tgcattcatc | cgagttcatt | tcgggcgtcc | ggtgaaattt | cacagcattc | 300 |
| acccttcatg | tgtagacaca | tattccacaa | attggttatg | atcaatgaac | tttttcaaag | 360 |
| acaatgtgga | aatcgtctct | tttcaaaagc | atgttggttt | tcagcttaac | aa | 412 |
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| -210- | 10205 | | | | | |
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| <211> | 387 | ĸ | | | | |
| <211> <212> | 387 DNA | ĸ | | | | |
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| <211> <212> <213> <400> tgtttgctta aagaccctct tcatggatgt ccacatctag gacgatcatc gacctccatg | 387 DNA Glycine max 19305 tgtctttgag cccgtcgact aaggggggtt cgatacatga tgacccata ccctgatacc | aagcgggaga agtgctccgg atgaacaacg gtctgcatca ggagaattcg aaatttagtc | aaggaaacat cctgtgatcc tctcttccca acatctcggg | caagatcgac cgttgctgat cgccttgaga cagcacaacc | gggacattcc agatatattc ggaaaccgag tgctacatgt | 120 180 240 300 360 |
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| <223> <400> | unsure at a 19306 | all n locati | ions | | | |
|-------------------------------------|--|-------------------|------------|------------|------------|-----|
| tgcgcatact | gcgtcaccaa | cacacagctg | gcttcgagac | ttcttagacg | atagaacaca | 60 |
| gcaccgctat | tctatgtnca | gcaccttgcg | caccatacat | gcgttacctg | tacctcaact | 120 |
| gtgaatcgag | ttcatacatc | cccctactat | cccattgctt | atgttgtata | catgctcctt | 180 |
| ctacaaactt | tgaattgcgc | ttattccatg | cccacgctta | ataggattac | taatgcacat | 240 |
| accttctcac | ccctcatgac | aaagcattgg | cttaaatcta | tgcctataca | cgaatctaaa | 300 |
| taacacctta | atgtgcttac | tgagatgtga | tactatgtca | ttgcccacat | gccaaagcct | 360 |
| aggcaaaact | cataatataa | ttgcaagata | attatactga | ccataggttg | tgacggtggg | 420 |
| tatatagggg | aaaacgcgct | taatcactct | caccgttctt | tcccaaccat | gcg | 473 |
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| tagctttgag | tgtgccgact | gaagccaaac | aagcaacgtt | tggtcatgca | ctcaggatta | 60 |
| acacgatcca | tgattcgatc | atcttctgtg | agaaattgac | gcggaatgac | tggattaacc | 120 |
| acaaacctct | gcaacttatg | cgatttaatg | actggttcaa | cgtgttgtcg | ccagtgtaga | 180 |
| tagttggaat | catccaattt | ctcagctatc | gttgtcggaa | atgaatgcga | actgaaacct | 240 |
| tgagatgacg | aagccatgta | tgtgaaggtg | cagctcgata | aactgaagag | aggagctttt | 300 |
| atgaagtaac | aagctntgat | accatatcan | aattagagaa | ctaagtgaac | aaagagaata | 360 |
| agagaaaaca | tt | | | | | 372 |
| <210> <211> <212> <213> <223> <400> | 19308 460 DNA Glycine max unsure at a 19308 | k all n locati | ions | | | |
| tagaacagta | tacttggcct | tcatttaact | gtctttggtg | tcttgttggc | cacgctcaac | 60 |
| aaagtacttt | cgacacctac | tgtacattga | tttcaccaat | gctgttatgg | gaatgttgcg | 120 |



| tc | | | | | | 362 |
|-------------------------|------------------------------------|---------------|------------|------------|------------|-----|
| <210> <211> <212> <213> | 19311 396 DNA Glycine max | · ‹ | | | | |
| <400> | 19311 | | | | | |
| agcttgaccc | ttacgagtca | gtttagtcaa | aggtaaagct | aacttggaga | aaccttctat | 60 |
| gaatetetge | gagtatcctg | ccaaacccag | aaaactccta | atctctaaaa | cagatttggg | 120 |
| actctcccac | tcaagaacga | cttctatctt | agatggatct | acagctatgc | ccccttgaga | 180 |
| tatcacatgc | cctaggaaac | taactttctc | taaccgaaac | tcacacttgg | acaacttagc | 240 |
| atagagttgt | cgatccctaa | gtgtatgcag | cacaatcctc | agatgttctt | catgttcctc | 300 |
| tctagtcttg | gagtatacca | aaatatcatc | tatgaatacc | accacaaaac | tatcaaggta | 360 |
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| aaagtctaac | cacttcaaga | ttcgtggcct | cttcaaatcc | cttgtttcct | gaaggaaatt | 180 |
| ccattcacag | accacccatt | ttcaatggtg | agggttacca | ttattggaaa | acccgtatgc | 240 |
| agatattcat | tgaagccata | gatctacata | tttgggaagc | aatagaaata | ggaccacaca | 300 |
| tacccactgt | agtagatgta | agcacaagca | ctacaaccca | taaacctaga | gataagtgga | 360 |
| cagaagaaga | taggagaaga | atccagtagt | caattacatg | ggatatcatc | ccaggaacaa | 420 |
| tttcag | | | | | | 426 |
| <210> <211> <212> <213> | 19313 408 DNA Glycine max | | | | | |

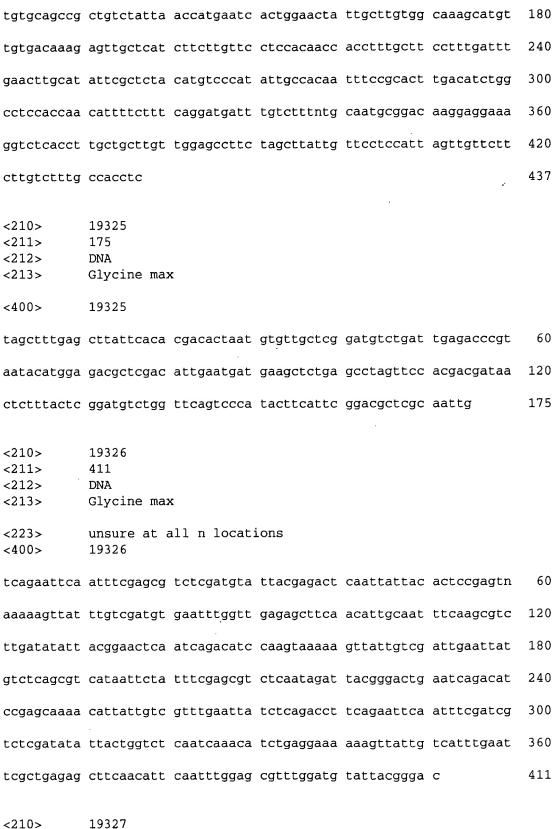
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|----------------------------------|---|--------------|------------|------------|------------|-----|
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| agttaaatta | cgaaaaaatt | ataatatatt | atttttttgt | aaaattagcc | atgagagttg | 120 |
| aaagatctaa | acgagaactt | tatatatcca | tctttatgag | tgaaatttct | tcaaatcttg | 180 |
| aațaaatgaa | caaaagaagt | aggcaatata | ataagctata | aagaaaggaa | caaacctcta | 240 |
| ttcttggaat | tggatgaccg | tgagttgtga | tttgtcttta | cttgtcacta | ttccaaatcg | 300 |
| taagcaataa | actgaatgtg | agaaataaag | agcagatttt | ggaagaaagc | aganatatgt | 360 |
| tggcaagcta | aacatgattt | aaggatatgg | atggcgtact | actagtcc | | 408 |
| <210> <211> <212> <213> | 19314 383 DNA Glycine max | | | | ı | |
| <223> <400> | unsure at a | all n locati | ions | | | |
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| atataaagaa | gaaaagagat | gggaaaaaag | agtacttgct | gacaaacagg | agctttcctc | 120 |
| attcttagtt | cgtctcacaa | caaaggtaag | atccttaata | tccttatcgg | ttacttgcgc | 180 |
| ccagtgtttt | aatttatcca | ttgctgaaat | gaatgcctgt | aaataaatat | atttgcatga | 240 |
| taccaatatg | tctgttccta | actcttttt | tttttttaat | tatcaatttg | atacccaata | 300 |
| tatattaaaa | gatcttggct | tgtgcgtgct | gtttggatgg | gttaaagtgg | tgatgatgga | 360 |
| gtttgagcag | tgacgttggg | ggc | | | | 383 |
| <210> <211> <212> <213> | 19315 414 DNA Glycine max 19315 | × | | | | |
| | | agcctctcga | tatgttacgg | gactcaatca | aacatccgag | 60 |
| | | aatttgctca | | | | 120 |
| | | agatatccga | • | | | 180 |
| _ | | - | | | • | |

| agaggttcaa | cattcatttt | cgagcgtctc | gatatgttat | gggactcaat | cagacatccc | 240 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| agtaaaaagc | tattgtcgtt | tgaatttggt | cagagattca | acattcaatc | tcgaacgtct | 300 |
| cgatatatta | cgggactcaa | tcagacatcc | gagtaagaag | ttattggtcg | ttgaattggc | 360 |
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| <210> <211> <212> <213> | 19316 422 DNA Glycine max | ς | | • | · | |
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| ctgagccaat | tcagacgaca | acaactttnt | gctcagatat | ctgattgtgt | ccagtaatat | 60 |
| aacgagacgc | tcgaaattga | atgttgaagc | tcttagcaaa | ttcaaacatc | attaagtatt | 120 |
| tactcggatg | tttgattttg | tcccgtcata | tatcgagacg | ctcgaaattg | aatgttgaac | 180 |
| ctttgagcca | attaaaacga | caataacttt | ttactcggat | gtctgattga | gtcccgtcat | 240 |
| atatcgagac | gctcgaaatt | gaatgttgaa | gctcagagcc | aattcaaacg | acaataactt | 300 |
| tctactcgga | tgtctgattg | agtcccgtaa | tatatcgaga | cgctcgagat | tgaatggtga | 360 |
| acctctgagc | caattcaaac | gacaataact | gtttactcag | atgtcggatg | ggtccgcgta | 420 |
| ta | | | | | | 422 |
| <210> <211> <212> <213> | 19317 333 DNA Glycine max | × | | · | | · |
| <223> <400> | unsure at a | all n locat: | ions | | | |
| agcttgaact | atatctagtg | agagtgtgaa | cttaaactgt | gagtgaacga | ctaactgtga | 60 |
| gtaatgatct | ttgcatgaat | ctctaaattt | tagaatgaaa | tgtataaatg | atgacatgat | 120 |
| gaaggccatg | attgtacata | cacaagctct | tttgaccaaa | tagcttacct | taaatgataa | 180 |
| ttgcatcctt | tgctcccttt | ttgagctgaa | tgatattgtc | aaaaaaaat | ttgaaccctg | 240 |
| aacttaaata | aatatctcct | gataccttgc | ttcgattcta | ggagagcata | tggtntaaga | 300 |
| caatttactc | taaatttggg | ggaggaaagt | caa | | | 333 |

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|-------------------------|------------------------------------|--------------|------------|------------|--------------------|-----|
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| tcagcttcaa | gctnttatct | tagattagga | atattgtgta | atggtccttt | gttgggcttt | 60 |
| tgtgatttct | tctattattg | tgtaactaga | accttgtcgg | acgaggaatg | gtgaattttc | 120 |
| tgtagagtaa | gcttatgagt | ctctcagtct | ctcaagcctt | ttaatgcaca | gcctactcac | 180 |
| ccagttttta | gtgatagtca | ctggtggaaa | ggagccaaga | gaattcaagg | atttctttgg | 240 |
| cgggtaagtc | acaaaggccg | cttacaaatg | ctagaagatt | gagattggga | ttatctaaaa | 300 |
| gtgatttgtg | cattgtgtgt | gaatcccata | gtgaaagtct | catatgacac | caattcggaa | 360 |
| ccatactttg | gttggtcaaa | acttggagct | gttcttttga | ttacatgatt | ggcttgaatg | 420 |
| gattcactat | aacttccaga | atgttgcc | | | | 448 |
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| <400> | 19319 | | | | | |
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| aatacgaaaa | atgatgaccc | tagggctgcc | aactcgtaaa | tcccgtgggt | atggcttttg | 120 |
| aaagggggga | aaagaggttt | ttgaatgcaa | aaacgtcccc | cctttcgtca | ttcttataat | 180 |
| tcgatgcacg | gatggctcgc | ccaggcgagc | taacctgcat | ttttttttg | agaggaacat | 240 |
| taaccatgtc | ccctccttcc | ttatgattta | acgtcttgct | taacttgaac | ttacttaagt | 300 |
| tagagttagg | cgttgattac | ttatttttaa | aacaaacaaa | tagtaagaca | actgcgaat a | 360 |
| caaaggat | | | | · | | 368 |
| | 19320 425 DNA Glycine max | | | | | |

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|-------------------------------|--|------------|------------|------------|------------|-----|
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| ggcacttctc | tctctttcga a | atttgctgag | gaaaattatt | tccgtgaaga | aaatccaagc | 120 |
| cgaggcgctt | ccgtaacgtt t | tccgtgagta | attatgcgaa | gattctcgac | cgttcttcaa | 180 |
| agattcatcg | ttcgttcttc (| gttttcttca | gtcttcaacg | ggtaagtacc | tcaaaccaag | 240 |
| cttttcaatt | cactctatct a | acccgtggtg | gtccacattn | tgtttcatgt | atttttattc | 300 |
| tcgttttcat | ttactttnta t | accccttt | tgacgtgctt | aagccgttta | tttaagtcat | 360 |
| ttctcgctta | atctaaaaaa t | aaaataaat | ttccaccgat | cgtttgaatc | gcatcatccc | 420 |
| gtaat | • | | | | | 425 |
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| <400> | 19321 | ~~~~ | | | | |
| | aaccaganag a | | | | | 60 |
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| cttcttgatc | ctctcacaag c | tcattggat | ataccccaca | ttcacttacg | aacgccgtgt | 180 |
| tgcaaaagca | acgcaggtat c | actatcaac | ggtgatgtta | ctttcacgta | gcacataata | 240 |
| ctgatctacc | atcattcaca t | aatatcaat | caacagcact | cctttcctac | ctcctattac | 300 |
| ctcccgctat | atcatcggaa t | caacggatc | acacatctac | angcgagacg | atgagaacga | 360 |
| atcattatgt | gaagaataac a | aaacagatt | cctaagtact | atgaaatcaa | tgccagaaag | 420 |
| acacacacct | tgtcatcgct t | ataccaatc | tgccaagtac | caaggccaag | cgtgattgac | 480 |
| caaatagata | tagegaceat e | tagatgctt | atcctctctt | acccagaaca | aataccgaca | 540 |
| caatactaat | gacagtaacg a | cactccgaa | acagagatcg | atcg | | 584 |
| <211> <212> | 19322 421 DNA Glycine max | | | | | |

| <400> | 19322 | | | | | |
|-------------------------------|------------------------------------|-------------------|------------|------------|------------|-------|
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| gcacccatat | acaatcaagg | cagcttcgtt | acctatatta | tttacacgta | cttccaaggt | 120 |
| gtatctgtta | cttacatcac | acacatctac | ttggctaaat | ttacatacat | gcatactcaa | 180 |
| agcatattgg | ggtacccaaa | attgcacatg | tgcacatcat | cgcatttcaa | atacctatac | • 240 |
| atacacaaac | ttcatgatga | atcttgacta | tctacacaat | aaggtgctac | attccatgct | 300 |
| tctttcaagt | cttcgctacc | taaacccgca | tgcaaattca | agcatattaa | cctttgctga | 360 |
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| g . | | | | | | 421 |
| <210> <211> <212> <213> | 19323 399 DNA Glycine max | | | | | |
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| agctttcctt | ttgctccctt | cctcacttat | accctcacat | gtttctccct | ctttaacatt | 60 |
| ttcttacttc | tggttaatag | atcttcttca | cgcaaagaga | tcgtaacctt | actctcacca | 120 |
| atttccagca | ccctaacacc | gctcattttt | actatgcatc | acatttgttt | tttttgttat | 180 |
| cttgtactta | ctcatggatg | taagggttac | tcatggatgt | aaggagtcag | gtaaattcgc | 240 |
| tatttaaatg | attataaccc | tccatcaatg | taattttggg | cctctactgg | cttattcctt | 300 |
| aacctttgta | tgcattaagg | ctcaaggaat | ccatgcatat | caccaagccc | taactgttga | 360 |
| cttatttaat | ccatatancc | taaatgtgga | aatattaat | | · | 399 |
| <210> <211> <212> <213> <223> | | : ill n locati | .ons | | | |
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| tttggaaaca | atggttttgt | caagttcttt | aaagaggttc | aggtcattgg | ttatatggtt | 120 |



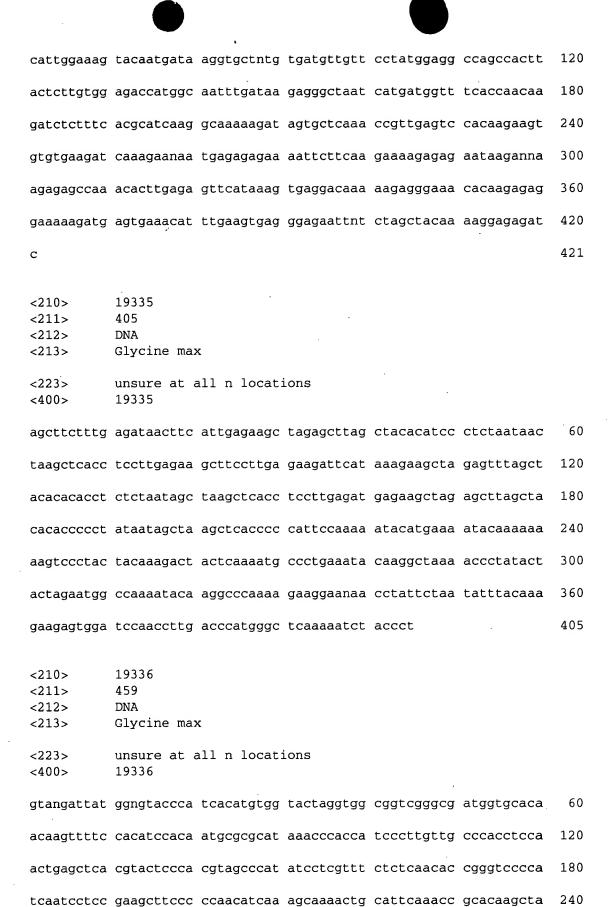
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|-------|-------------|--|
| <211> | 380 | |
| <212> | DNA | |
| .212. | Claraine ma | |

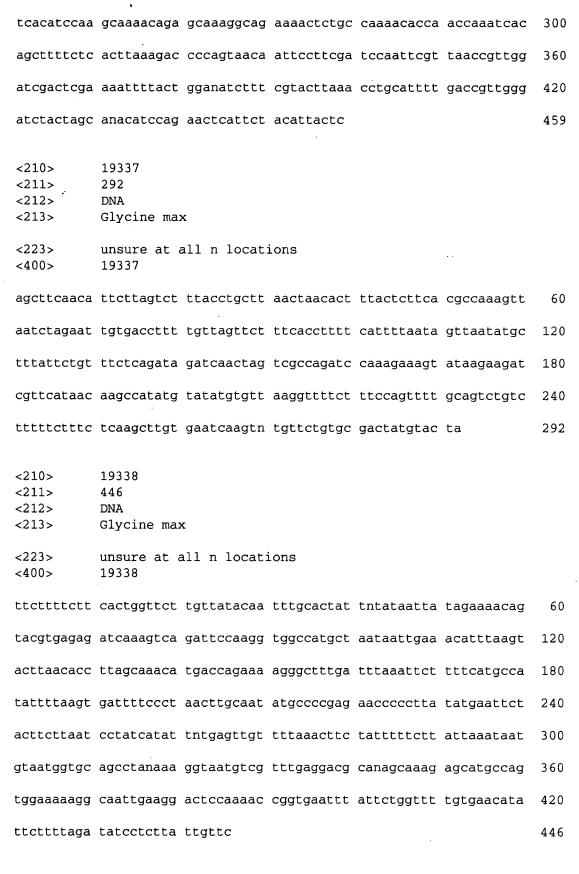
<213> Glycine max

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| aatttagttt | ctgttataat | tttaattaaa | atatttaatt | acataattag | tgaaaaacac | 120 |
| ggagctataa | tcttttcatc | atgattttct | ttaatctttg | atatttttt | ttcaattttt | 180 |
| atgcttttca | ttattattca | gtctatccta | tttttcttta | tttaatatçt | tccttgataa | 240 |
| tcttacatcc | tttataattt | atactctatc | atttttatat | ctaaatagaa | tcataataat | 300 |
| ttgtttaaca | aaaaaattct | tatattataa | ctcacaagct | antatgtgaa | agatgaatac | 360 |
| aaaatataat | gtatgtattc | | | , | | 380 |
| <210> | 19328 | | | | | |
| <211> <212> | 205 DNA | | | | | |
| <212> | Glycine max | ζ | | | • | |
| | _ | | | | | |
| <223> <400> | unsure at a 19328 | all n locati | lons | | | |
| <400> | 19328 | | | | | |
| ntaattaagt | gtatcacttt | tgttattatc | atcatgcctt | tgttttgtta | acttgaagtt | 60 |
| atttttttt | tttaaatacc | tgaggtcaat | atccaagata | tactaaatta | ctaatataga | 120 |
| tatacccaaa | cacacatgaa | gctatctgat | ttgttcactc | agtttgtccc | ccctgttatg | 180 |
| gatgcataat | agagggggg | agggg | | | | 205 |
| | | | | | | |
| <210> <211> | 19329 392 | | | | | |
| <211> | DNA | | | | | |
| <213> | Glycine max | ζ | | | | |
| <400> | 19329 | | | | | |
| agcttcctct | ttaagcttct | tatccaagcc | actctcttgg | tggtgaagct | tctccttcca | 60 |
| tgacttattc | tctagcggat | gacgtctcct | ctaacctctt | ctcctttatc | tttcgctgca | 120 |
| attccatggc | taataatcac | cattgaagga | ccttattgaa | gctcaaagat | ccagcctcca | 180 |
| tagaagcttc | ataagcaagc | ttccaacaag | tggtatcaga | gcacaagagc | ttcaagtagg | 240 |
| tgctccttaa | acctccacta | attttcagct | ttactttctc | ctccattgtt | gttgcttcgt | 300 |
| ttctctccat | gtatctcctc | acgtgtcttg | tgctgaatgt | tgtaacataa | tttttagaag | 360 |
| | | | | | | |

| ttccaccgat | tagcttgcta taa | ıagctaga | tt | | | 392 |
|----------------------------------|------------------------------------|----------|------------|------------|------------|-----|
| <210> <211> <212> <213> | 19330 394 DNA Glycine max | | | | | |
| <400> | 19330 | • | | | | |
| tgtcaaagct | gacaatatct tca | ıgatatgt | ggacaccggg | aatctaatac | tgacttccca | 60 |
| gtaggttcaa | acgtgactca ggt | gtttgtt | gatcaactac | cgtgactaaa | cacccttggc | 120 |
| atagctttgg | ctcgcataga ttt | tgcacca | tagggtttga | acgctcccca | cactcaccct | 180 |
| cgcggcactg | agatccttat agt | ccttgag | ggtactcttt | atgttggatt | tgtgacttcc | 240 |
| aatcaagatg | gaaatcgcct ctt | caccaaa | gtgctgaaca | agggtgatgt | gtttgtgttc | 300 |
| ccaattggtc | tgattcattt cca | aatgaat | atgggaaatg | ggaatgctgt | tgccattgct | 360 |
| ggccttagca | gtcaaaatcc agg | agctatc | acta | | | 394 |
| <210> <211> <212> <213> | 19331 398 DNA Glycine max | | | | | |
| <223> <400> | unsure at all 19331 | n locati | ions | | | |
| agcttgtagt | ttattcgaac gac | aataaca | tttcactcgg | aagtccgatt | gagtcccgta | 60 |
| atatatcgag | acgctcgaac ttt | aaaaccg | aagctcgtag | cagatttgaa | cgacaatgac | 120 |
| atttcactcg | gaagtcctat tga | gtcccgt | aatatatcga | gacgctcgaa | atttagaatc | 180 |
| gaagctcgta | gaaaatacga aca | acagtaa | cttttcactc | ggaagtccga | ttgagtcccg | 240 |
| taatatatcg | agacactcaa aat | ttaaaac | ccaagetete | aganacttct | aacgacaata | 300 |
| acttttcact | cggaaggccg att | gagtccc | gtaatatatc | gagacgctcg | aaatttaaaa | 360 |
| ccgaagctcg | tagcaaattc gaa | cgacaat | aacatttc | | | 398 |
| <210> <211> <212> <213> | 19332 468 DNA Glycine max | | * | | | |

| <223> <400> | unsure at a 19332 | ll n locati | ions | | | |
|-------------------------|------------------------------------|-------------|------------|------------|------------|-----|
| ntggttntaa | atttcgagcg | tctcgatata | ttacgggact | caatcggtct | tccgagtgaa | 60 |
| aagttattgt | cgttagaatt | agctgcgagc | ttcggtttta | aatttcgagc | gtctcgatat | 120 |
| attacgggac | tcaatcggac | ttccgagtga | aatgttattg | tcgttcgaat | ttgctacgag | 180 |
| cttcggtttt | aaatttcgag | cgtctcgata | tgttacggga | ctcagtcgga | cttccgagtg | 240 |
| aaatgttatt | gtcgttagca | tttgctgťga | gcttcggttt | taaaattcga | gcgtcacgat | 300 |
| atattacggg | actcaatcag | acttccgagt | gaaatgttat | tgtcgttagc | atatgctgcg | 360 |
| agcttcggta | ttaatatttg | agcgtcttga | tatattacga | ggactcatcg | gacttccgag | 420 |
| tgaaatgtat | tgtcggtcaa | attgctcgag | cttcggttta | attcgagc | | 468 |
| <210> <211> <212> <213> | 19333 432 DNA Glycine max | · | | | | |
| <400> | 19333 | | | | | |
| agctttcaat | gtttataaaa | accaaaaaac | tttggaaagc | ttttggcaaa | aggaagaaga | 60 |
| agaagaagtt | caaagagact | cagaaatcaa | tgtggaaaac | ttgcttgtga | aaagaatgaa | 120 |
| ttggaaaaga | ttgattgata (| gaatgaatga | atgaaaatgc | aaaacaaagt | cttgctttta | 180 |
| tagactcttc | atgtcttgtc a | aagaagacca | tttagaagag | ttataaattt | tagaaaaact | 240 |
| taaaactaat | ttgaaaaagt | caaaaacctt | ttgaatagtt | acatcttttg | atttattcag | 300 |
| aaacaatcac | tggtaatcga | ttaccaaatc | actgtaatcg | attacacaag | gcttttatgt | 360 |
| gaaaggatgt | gactcttcac a | attcgaattt | gaatttcaat | gttcaaaggc | actggtaatc | 420 |
| gattacaaaa | ac | | | | | 432 |
| | 19334 421 DNA Glycine max | | | | | |
| <223> <400> | unsure at a 19334 | 11 n locati | ons. | | | |
| ntaatggctt | agtgaggatg g | gagaggtgca | agtaaggaag | caagtagagt | tggatatttc | 60 |





<210> 19339

| • | | | | | | |
|----------------|-------------------|-------------|------------|------------|------------|-----|
| <211> | 397 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | K | | • | | |
| <400> | 19339 | | | | | |
| agcttgtccg | ttggatgcct | acccattacc | cattagaaat | agactagtag | atagggcagc | 60 |
| aagaccctgc | ctacttagct | tcttagatgc | atactcaggg | tacaaccaaa | tacggatgca | 120 |
| tccacaagat | gaggagaaaa | caaacttcat | aacctagtcg | tctaactatt | gctatcagat | 180 |
| tatgccattc | ggcctaaaaa | aggctagctc | cacttaccag | cacctaatgg | acatgatatt | 240 |
| caaagaacaa | attggaaaga | aaatggaggt | atatgttgac | aacatggtgg | taaagtctaa | 300 |
| tgatgcagaa | tcacacacct | atgactcgga | agatatattt | gcaaagatct | gaaagcataa | 360 |
| catgtaactc | aattcgaaga | agtgtatgtt | tggggta | | | 397 |
| <210> | 19340 | | | | | |
| <211> | 158 | | • | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | • | | | | |
| 12207 | 01701110 11101 | • | | | | |
| <223> | unsure at a | ll n locati | ions | | | |
| <400> | 19340 | • | | | | |
| gttgaacaat | gaactattct | tggaagagaa | agtattgaac | attttctctc | tataaagatg | 60 |
| aacaacaatg | aatgatcaag | ttctgtatgt | ctcgntttgt | aaacaagttg | ttattgaaat | 120 |
| tcatgccatg | gtcacatatt | tatactcaat | tgatgact | | | 158 |
| <210> | 19341 | | | | | |
| <211> | 400 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | : | | - | | |
| .222 | | 7.7 | | | | |
| <223> <400> | unsure at a 19341 | II n locati | ons | | | |
| (400) | 19341 | | | | | |
| agcttttcga | ttcattctat | gtacccgtag | tggtccacat | tgtgtttcgt | gcattattat | 60 |
| tctcgttttg | tttacttttt | ataccccctc | ttgacgtgct | taagccattt | tacttaagtc | 120 |
| atttctcgct | taacttaaaa | ataaaataaa | ttttcaccga | acgtttgaat | tgtattatcc | 180 |
| gttaacttcg | gttaaaataa | attccgaccg | ttcggtcatg | ccgtaaccac | gttggaaatc | 240 |
| aaaangaggt | cttctaatac | atatatatct | atctcncnnn | gacatctttt | agtaaaataa | 300 |

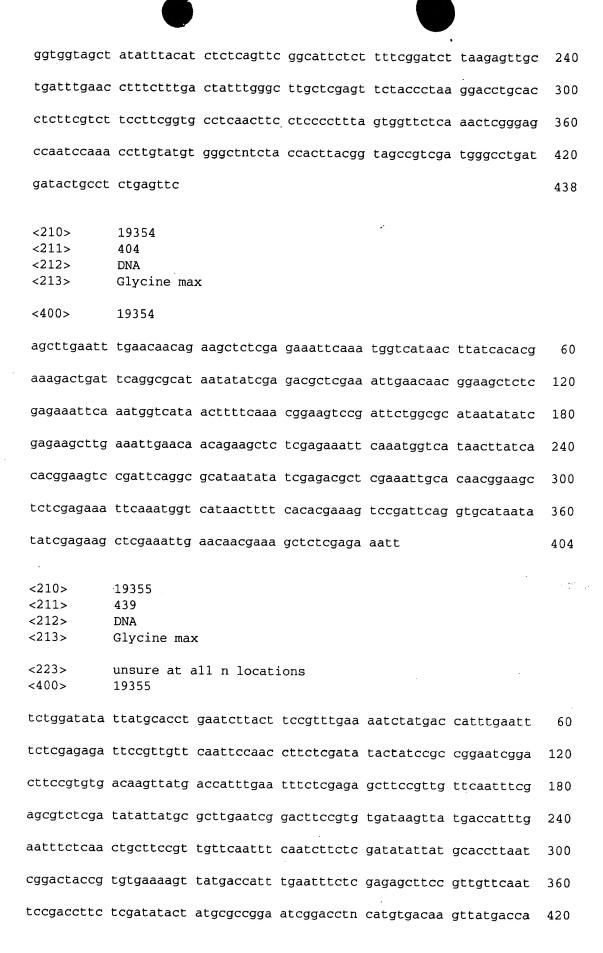
| agcggaaaat | caatcggaca | ttttctcttt | gggatttctc | attcttaatc | gaattgatta | 360 |
|-------------------------|------------------------------------|------------|------------|------------|------------|-----|
| ataactaaag | tgaaactaat | gctaaaatca | actcgcctag | | | 400 |
| <210> <211> <212> <213> | 19342 382 DNA Glycine max | × | | , | | |
| <400> | 19342 | | | | | |
| agcttgttat | atgcacaatt | aaggtacaca | cttttgatat | tcttttgctc | agacttggca | 60 |
| cactattcca | gagcgcctat | ttcgggaaaa | aagccctaga | agcagtaaga | ggagcagctt | 120 |
| gagcattgaa | gcctatgttt | tggaatttgt | gagagattac | tgagctggat | agtgagtgtg | 180 |
| agatgttgag | aagaggagga | cgaatccccc | atcttgtgta | aggaactatc | attctctgct | 240 |
| tttagtctca | tttattgata | cggttgcttt | gtaatggctg | gctaaacacc | ctagttgggg | 300 |
| atctatgatg | aacaactgat | gtcaatactt | actatctaac | tgagtatata | ttgtgagtgc | 360 |
| agtgcttcct | ctcatgctta | at | | | | 382 |
| <210> <211> <212> <213> | 19343 463 DNA Glycine max | × | | | | |
| <400> | 19343 | | | | | |
| tgagctatca | gaagacttgc | ttattcattt | agttctaatt | tatctacctt | catagcttag | 60 |
| tcagcttaag | atctcttata | acggtcggaa | ggagaaatgg | tctcttaatg | agttcatttc | 120 |
| atactgcgtg | caagaagagg | aaagactaaa | gtaagaaacg | actgaaagtg | ctcatgttgt | 180 |
| gagtacctct | aaagacaagg | gcaaaagata | aaggactgat | gagcccaaca | atgaagctgc | 240 |
| ttagggtcta | ggacaaaata | taaaaaatca | aggtgacaac | tatttctttt | gcagtaagcc | 300 |
| tggacatgtc | tagaagcaat | gtaaatgtac | catatatcat | gcttggcgtg | caaagagagg | 360 |
| tatgtttctt | actatggtat | gatctgaggt | caatttacct | tcagtaccta | caaacacttg | 420 |
| gcggtagact | ctggtgccct | actaacatca | gcgtttcaat | gca | | 463 |
| | • | | | | | |

| <213> | Glycine max | | | | | |
|-------------------------------------|---|------------------|------------|------------|------------|-----|
| <400> | 19344 | | | | | |
| gcctcgctcg | tttcattatt cc | tagctata | atagagtgct | gcactgaaac | atattctatc | 60 |
| tatctatgca | gagtaatgcc cat | :gaccata | gccctataac | gacttgtggg | aatgcttgga | 120 |
| ctctacctag | tacctacage aca | actgccta | ctgacctgct | aaccatataa | agcagacctt | 180 |
| ccattggtcg | aacataccat gca | a | | | | 203 |
| <210> <211> <212> <213> <223> <400> | 19345 410 DNA Glycine max unsure at all | n locat: | ions | | | |
| | agcetgetee eta | anneanna | ccatcaacca | tctagcaatt | cttcattcca | 60 |
| | ctttctccct ttc | | | _ | | 120 |
| | | | | | | |
| ctttgatttt | ccttttatct gca | acccctgg | cttctgtatt | tggacctctg | gtttgtgttc | 180 |
| tctgttaact | gccccttgtt cct | cagcttt | tgtaccagta | ttacgtatgg | tggtataaaa | 240 |
| attacaacaa | agggagcata ato | gtagaatg | ttcttattat | gtcgtagttt | tgaactgcat | 300 |
| aaagagcttt | cntaagagtg tat | attatgc | taatgaaaaa | cgatacgttc | actgcacatt | 360 |
| caaaaatcat | ttatgttaat cto | ctaagtat | gcattataat | attcataatt | · | 410 |
| <210> <211> <212> <213> | 19346 380 DNA Glycine max | | | | | |
| <223> <400> | unsure at all 19346 | n locati | lons | | · | |
| ttatcttaga | tcttccatgc cca | agctgata | atgtcaatcc | tgatactgct | agaggatctg | 60 |
| gtccccacaa | agacaaattt cad | cagttacc | taggggtggt | ggctagagat | gaaataccaa | 120 |
| ttgtccactc | caattggaat ctt | :gt <u>ctcgg</u> | acaatctaaa | gaacctaatt | tacgaagaca | 180 |
| ttttggtaag | tccttttaat acq | ggtgtatc | tttatcattg | cctttataac | acatttttac | 240 |
| atcactaatt | aagttggtgt tat | ttaacac | atttgttgta | tagcagaaat | ntgacatccc | 300 |

| tgaacgtgac | aatgcgaaac | aaaaaggtca | tgtctatagt | ggcaactaca | tggaggcaat | 360 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| ttaagtcttc | gttgacatct | | | | | 380 |
| <210> <211> <212> <213> | 19347 475 DNA Glycine max | × | | | | |
| <223> <400> | unsure at a | all n locat: | ions | | | |
| ntgtctttct | tagtactcat | aattctacat | taattatttt | tcttgatgca | tgattaacaa | 60 |
| aactttaaca | tgcagaattg | ataagaggta | aatgtcattg | caacaatact | atgtggtaca | 120 |
| ttcttagtat | gatcacccct | gcagaaccat | aaagattgtg | aataacaagt | caacatcact | 180 |
| tagtgccatg | gcaatggtaa | agccaattac | gccatttcct | taggcctccc | acaaagctct | 240 |
| taattctata | tatatttata | aaaaaataaa | tcaaatttaa | attatcaggt | aatatattct | 300 |
| attgcattat | ctattattat | ttgcgtagcc | agattataaa | attttaatta | cacatagata | 360 |
| tattataaat | catttgagaa | gtttataatt | catttgacaa | ttatagaaaa | attagtttcc | 420 |
| atctacttta | aatcttgtaa | aatcacactt | aatgaatata | atgattcaat | ttata | 475 |
| <210> <211> <212> <213> | 19348 383 DNA Glycine max | ĸ | | | | |
| <223> <400> | unsure at a | all n locat: | ions | | | |
| tagcttggtg | atctttgctg | ctggttaaca | ctgagcttgc | ccaccctacc | attactaagc | 60 |
| atttcactgc | taatagaaga | tgcaatggat | agctttttct | tgcatggtga | atagctatca | 120 |
| aagttcatgt | ccacattccc | cccagtcctt | atttttgggg | agcttgttga | tcttgctttt | 180 |
| gcttttgctg | attctgttgc | aaccatgtat | gctgganttg | catgatagct | taattatgag | 240 |
| tggtcctctc | ccactgaaca | ctgcctccta | tgaggaaaag | atcttcttga | accaagtatt | 300 |
| ggagaatcca | atccttcaac | tggattttgc | ctctgaatat | tacttctcag | cttaagttgc | 360 |
| cttcctccat | attcctcccc | aac | | | | 383 |
| <210> | 19349 | | | | | |

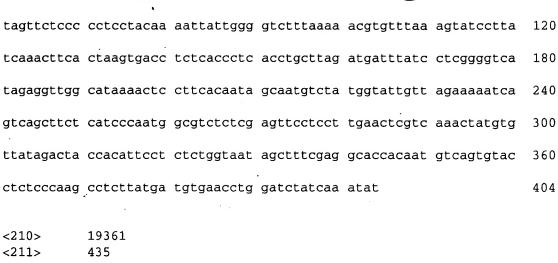
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|----------------------------------|--|-----|
| <223> <400> | unsure at all n locations 19349 | |
| tagcattagc | atgtgttgac actggatcac taatcattgt gtacgtgttt gttactttag | 60 |
| tggatcctat | caaagaacct ccgcttataa ctacaaacat agttctgtct gtttttgttg | 120 |
| tgtattgtca | tgtggacaaa gttccatgcc atgtttgaga catctaagat tggcgtcttg | 180 |
| cctttgccca | gtattatttt ttgcaacatc ttcctttctt aaccttgtca ttacctcgaa | 240 |
| attntaatat | ggcaacttac tcttgtggaa aataattttt aagaattaat ataacacttt | 300 |
| aaaattaaat | ttagaatatt aaaaaaatat aaaacataga tataattctt taggtgctat | 360 |
| tgatactttt | ctcctgttag aattaaaatc gtacttcagt aatccacata ataaagatat | 420 |
| acatcataaa | attacatcaa ttatcatagt gaaactct | 458 |
| <210> <211> <212> <213> | 19350 425 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19350 | |
| agcttttcaa | ttaatctttg gctagctaca ttagtgcaat tacccctatc aataatcaaa | 60 |
| gagcatattt | tccccatgac catgcaccta gaatgaaaaa tgttctccct ttaagtttca | 120 |
| tctctatcct | tacacacatt acccattaac ctcctaacca taaaaagatc accttccagg | 180 |
| ggttgtacat | cacattcatt ttcactctca ctagaagaac tagaccagct agaagaagat | 240 |
| aaactaatga | tatccccatt acccaacaca accatagtcc ttttgctagg acattgngag | 300 |
| gcattatgac | cctttcccaa acacttanaa catttaatag aacttacttt tgaagaagta | 360 |
| ggagtagggt | tagaaccaca ccatactacg agagaggttc cctcttttac catttttaat | 420 |
| atccc | | 425 |
| <210> <211> <212> <213> | 19351 480 DNA Glycine max | |

| <223> <400> | unsure at a 19351 | all n locat: | ions | | | • |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| tcaatgatnt | aacaatngac | tnggtagaat | tgcccacgtc | atggttgtca | tgcactgccg | 60 |
| cattgccacg | tggaatgttg | aaacttgaaa | tttattatga | aaatattgta | ccctataata | 120 |
| acagcgtatg | ctaattgaat | tgtttaaaat | taaattttga | ttgtttatat | aaaaaattat | 180 |
| cattacaaat | ttttaaattt | aaatccattc | aaaataaaaa | taaaaaatta | ctcaaaaaac | 240 |
| taaaatttaa | aataaattat | ttctttttta | tttttaaaat | caaaccaaat | cgtaacactt | 300 |
| atattaatat | ttattattt | aattntatgt | gtgtattatt | ntaattttat | aatattntta | 360 |
| tgatattatg | tatatgaaat | ttatataata | tatattttt | tcaaataact | ttctaaagat | 420 |
| atanntttat | ttanatagat | gaaaatttac | atatctntat | catanaaggg | ttaatatatt | 480 |
| <210> <211> <212> <213> | 19352 309 DNA Glycine max | ĸ | | : <i>,</i> | | |
| <400> | 19352 | | | | | |
| agtttcatgt | ttattcaaga | ttgattcaaa | gaagttctaa | tgattacaaa | tgtgatgaca | 60 |
| aaaagctcac | aggttaataa | cacttcatga | taacaaagat | gatgatctca | agaatcaaag | 120 |
| aatgagttca | agatgttcaa | gattgaatca | agaacatttc | aaggttcaag | aggaaatttg | 180 |
| atttctagat | tcaagaatca | agagaagact | tagtcaagat | aagtatgaaa | acatgttttc | 240 |
| aaaaactgag | taacacatgg | attattctca | aaacctggtt | accaaagagt | ttttactctc | 300 |
| tggtaatcg | | | | | | 309 |
| <210> <211> <212> <213> | 19353 438 DNA Glycine max | ĸ | | | | |
| <223> <400> | unsure at a | all n locati | ions | | | |
| tctcttagac | cttaggcaaa | ccttcaactc | atcctttttt | atttttctgt | ctacttgaga | 60 |
| taggtccatt | tcctctctcc | ggagcttaaa | ctcgctgcta | ctgccccaca | aagcccctcg | 120 |
| gaatttgttt | cggccatgtt | cttccctacg | agcccttttg | gtctcttgtt | ccaaggcctt | 180 |



| tttgaagttc | tcgagagct . | | | | | 439 |
|----------------------------------|------------------------------------|---------------|------------|------------|------------|-----|
| <210> <211> <212> <213> | 19356 382 DNA Glycine ma | · x | | | | |
| <223> <400> | | all n locat | ions | | | |
| agcttcatac | tatgctttat | taatggnggg | cggaaaggca | aatgtcagtg | cctgctcaac | 60 |
| aattaatcca | tttgatttct | catggaaaga | tctatctaca | tatattgtaa | ataatatctc | 120 |
| ccgttataga | ttttctaagg | ttaattcaac | aatggaatca | gctagcgcat | gtgcattaaa | 180 |
| aaggatttga | tgctctgaaa | cctctattgc | ccaagaaacc | atccttctcg | ccaaatcaag | 240 |
| tcggtgaaga | atttgtcgaa | tggactaatc | aattcacacc | attatccggt | gtgattggaa | 300 |
| atatagtcgt | aagcgtcttg | ttgccattac | taaagcanaa | actaccttct | ctaatgatct | 360 |
| ttgaagttcg | accccctgaa | gt | | | | 382 |
| <210> <211> <212> <213> | 19357 442 DNA Glycine max | ĸ | | | | |
| <400> | 19357 | | | | | |
| tttctttcac | aatcaatttg | tctactgact | aacaatttta | aatgcatgtt | cacattettg | 60 |
| ttctttctta | gtctaacata | cacacttgct | caaactcatg | ataagaaaca | caaactccat | 120 |
| cacaatcatg | cacttaattt | aaaataaaag | catataacta | ttttcacaaa | aagataaaaa | 180 |
| gtgttttact | accatgtcat | caaaaacaag | tcaaactatt | caaaatgctt | caggataagc | 240 |
| aaactaacta | cccataaata | aaactagcag | tgtatgtaga | cctaaaggaa | atattgtatg | 300 |
| aaaaccaaaa | ttgtaataat | aataataaat | caaaaagcaa | aaagtattat | caggaatcaa | 360 |
| aattcatgtg | actggtcttg | gatatcctgt | gcctgaacat | cctccttatc | tgtcaaatgc | 420 |
| aatactggag | tagtcggagg | ag | | | | 442 |
| | 19358 391 DNA | | | | | |

| <213> | Glycine max | |
|-------------------------------|--|-----|
| <223> <400> | unsure at all n locations 19358 | |
| agcttgaga | a tacagaatgt tatggttgaa ataagcaaga taaatattgg aagtttatct | 60 |
| agataatta | c ctgacttaca tgtgtgattc taaatctaat cattatatgt agaagcttat | 120 |
| ctcagataa | a ccttcaatct ctttttatgt tctttagtga ggaactggta aagattcaat | 180 |
| gctttgatad | c aatctggtga gtgagtgtcc gagggtagat aattactgcc atgtatttat | 240 |
| gttactcttg | g catatacacc actacactca gtctaataca cccttagctt anggtacaat | 300 |
| anaagagtga | a ttcctagttc cttagttttt cgaanatatc ttanattagc actaggaaaa | 360 |
| anatactgta | tggagatett catatettat g | 391 |
| <210> <211> <212> <213> <223> | 19359 433 DNA Glycine max unsure at all n locations | |
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| | ctaacaattt ccattaatca atttattatt atttacttgt ccttagattg | 60 |
| | aataagatca atgctttgag gaattatttt atatttcata atgtggaagc | 120 |
| | aattaaaatt ttattttgat gtgaattata gcaatgggat caacacaaat | 180 |
| | aggacaagag aattagtcat ttacattgca tgataaagaa aactagataa | 240 |
| | aaaatgctgt tgataatacg aaagttcacg agaccagtgc tactaatagt | 300 |
| | gtgtatgttt ttccctaaac cctaaagaca cataagatgt gtaacaaacc | 360 |
| | agtatagaac aagcagggct ggtttaacag catcaaataa tctatcanta | 420 |
| aaatgctatt | gct | 433 |
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| agcttatctt | tcaactcctg aggattaggc ctcaagagtg caaatatgga taatgtacac | 60 |

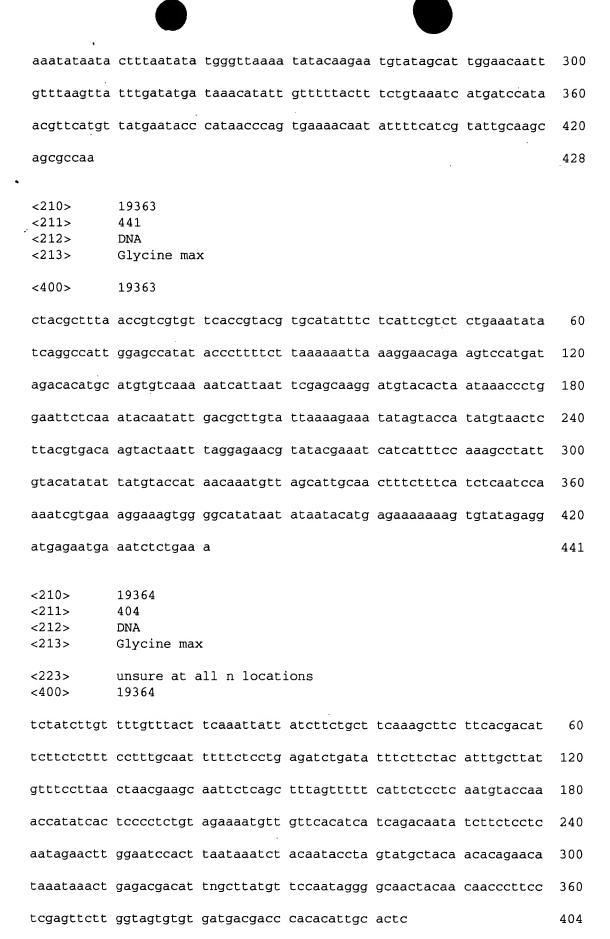


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| <213> | Glycine max |
| | |
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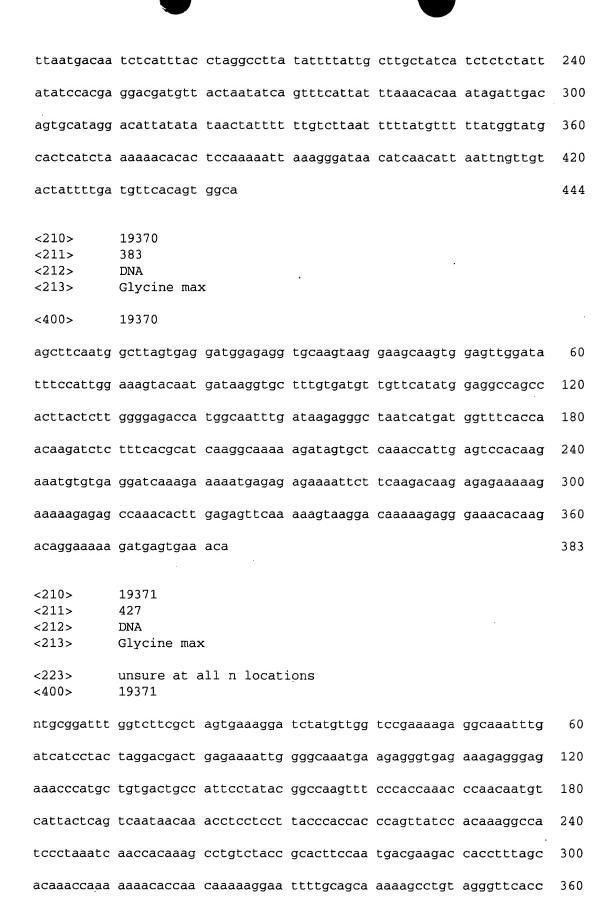
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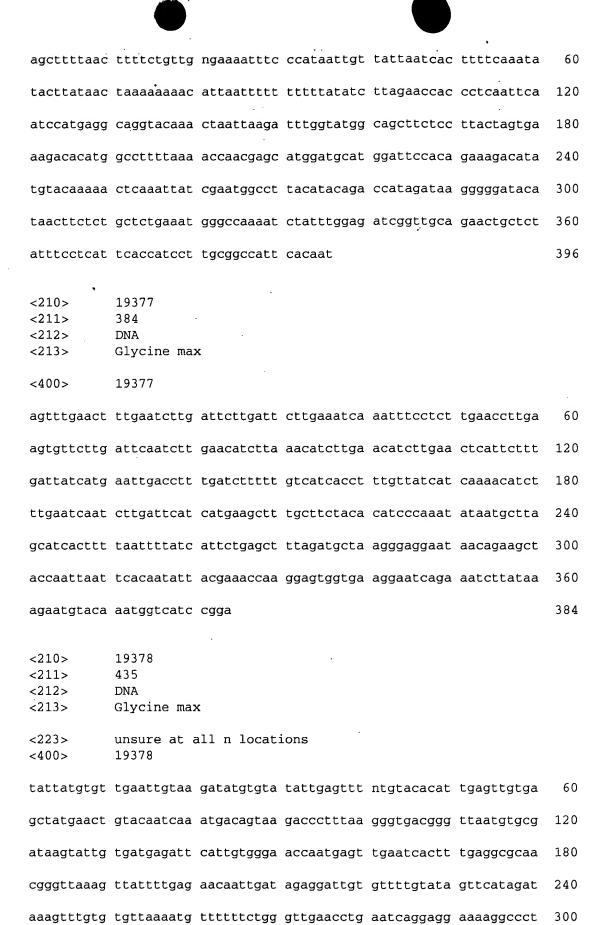
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| ttccctcact | cctacaatat | ttctacatca | tccacacttc | ctatatccca | aaagccaaaa | 60 |
| aaaaaaacat | caaaaccttg | aaagagaaca | cttttatgaa | actttggaaa | gtgcacatag | 120 |
| aagaaaacaa | aattggaaga | taagagggga | aagaggaggg | cctcttacct | ctaaaatcaa | 180 |
| tccaaattaa | aaaaaaaag | aaaaaagaa | aaaaagctca | acaacaaggt | cctttgtgtg | 240 |
| agggtttcca | tgtcttgcac | tctctaatga | tggattgtct | cagagaggaa | gaagagagtg | 300 |
| aaatgagagt | attttgtgtt | gtggttcagt | ccanacactc | atctcaatcc | aaacactcta | 360 |
| aaacttatta | tccacacctt | anacaaggtt | ttcgcataca | taattgtaat | catatataan | 420 |
| aacatcacaa | gtcatca | | | | | 437 |
| <210> <211> <212> <213> | 19366 400 DNA Glycine max | ĸ | | | | |
| | 19366 | | | | | |
| agctttaaaa | ttgtaacttt | ttaaattaga | aaatatgatt | aaagataact | ttaaaagttt | 60 |
| gaaccctaaa | atctttttag | ttacttttaa | cactttttt | tataaaacct | atgcttaaga | 120 |
| aaaaataata | agaaaatttt | aaaccctaaa | ccccaaaaga | tttttttt | ataaaaagtg | 180 |
| tgaaagtaac | aaaaaatatt | ttagggttta | aacttctaaa | tgtatcttta | atcatatttt | 240 |
| gtaatttaaa | aagttacaat | tttaaaattt | ttaaaaaaat | agtcaaggac | taatttgaaa | 300 |
| aaataaaaaa | taattaatga | gaagatgaca | tataatttga | tagattaaaa | aattaccaat | 360 |
| atcataaccc | ctaatattat | tggaaacacc | atagaaactc | | | 400 |
| <210> <211> <212> <213> | 19367 436 DNA Glycine max | c | | | | |

| <223> <400> | unsure at al 19367 | ll n locati | ions | • | | |
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| tgatcaaaac | aattctctaa c | ccattccaat | cctttcttat | catacaattg | cttattcaaa | 60 |
| tcattctcaa | acactcattt c | cataccaaat | aatccactgc | atatcatttt | caatcaattc | 120 |
| attgttcaaa | cacgcttttg g | gtacaagtaa | acaactcaaa | gtgttgaaat | tttaaataaa | 180 |
| tgaaatataa | aataactgaa a | cataaaaac | tgaaattaaa | ataactgaac | ataaatcata | 240 |
| aaataattga | aaataaacta a | aatgttcaa | gatgcacaaa | tttaaatgtc | atgttcatca | 300 |
| tgtggctagt | cttcattaag a | tccagtgct | ggagctacta | atgaatcctg | gataggttgc | 360 |
| tttggctccg | tgactggtgt a | ıgatggctgg | gtctcctcac | gaggaggtgc | agaggatggc | 420 |
| tcangtatct | gatctg | | | | | 436 |
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| agcttgattt | ataaaactgc a | gttttattt | ttaaaaaaat | tatgaggcga | agttagaaaa | 60 |
| aaaaaaaggg | ttgagagtaa c | catatcttta | taattgtccc | gtcccaccgt | aatagagaaa | 120 |
| attttcgtag | tgaaattaaa t | atgatctca | acaaccgtgc | ataattaata | tatcattttt | 180 |
| ctcattagaa | ttgaacagaa a | ataatgatg | ttatttgtta | ctttctaaat | tgagggtata | 240 |
| atagcatcaa | tgcctaagat a | aacggtgac | aatatcaaat | ctatcaatca | atcaaaangt | 300 |
| taaagggtat | gtattttcta c | ctccat | | | | 326 |
| <210> <211> <212> <213> | 19369 444 DNA Glycine max | | | | | |
| <223> <400> | unsure at al 19369 | l n locati | ons. | | | |
| | | | | | | |
| gacactatga ' | aactccgctg c | tcaaggtgc | cátgcttttt | ttaattcctt | ttactaatat | 60 |
| • | aactccgctg c | | | | | 60 120 |



| ccaaattccg | ttgtcatatg | ctaaacttga | tcccatatcc | actcaataat | tcaatggtag | 420 |
|--|--|---|---|--|---|---------------------------------|
| ccataac | | | | | | 427 |
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| gttgcatgta | agctgggtgc | tattacgata | gccgactact | tggcggacac | tatggcctgt | 60 |
| tatcatcata | tgcagtagaa | atattagtct | tgtatattat | caatcgctct | cattcagtag | 120 |
| tgcgtggtcc | tctagaggtg | agtcacatat | tccatcagta | atatttttgc | aacctttttt | 180 |
| ttaatatact | tacagaataa | aaccattgaa | ctgctactac | tagcaggtgc | tatacatatt | 240 |
| tgtggactac | tacagctcat | ttgattgcga | ccataattat | gttagtatat | ggggtccaaa | 300 |
| atccttatcc | tctcttccag | aaattgttgg | taagttatgc | aatgtgtaag | caactgcaga | 360 |
| ttttcaatct | tcagtccttg | atatcccaac | tcccttgc | | | 398 |
| <210> <211> <212> <213> | 19373 432 DNA Glycine max | × | | | | |
| <223> <400> | unsure at a | .11 - 1 | | | | |
| | 19373 | all n locati | ions | | | |
| tctgtttcaa | 19373 | | | agtgaataaa | ctgttgtttt | 60 |
| | 19373 cgaaatgggt | aacaggactg | attatattta | agtgaataaa taaagattta | | 60 |
| gtaatttgaa | 19373 cgaaatgggt gccacacaat | aacaggactg acttcaaaag | attatattta ggtaaatgaa | | tcaccccttt | |
| gtaatttgaa tgaatcactt | 19373 cgaaatgggt gccacacaat tcatcgtccc | aacaggactg acttcaaaag tttttgacct | attatattta ggtaaatgaa tacatatttg | taaagattta | tcaccccttt tacgtacgct | 120 |
| gtaatttgaa tgaatcactt gcagcctgtt | 19373 cgaaatgggt gccacacaat tcatcgtccc aacagttgct | aacaggactg acttcaaaag tttttgacct acttttaagc | attatattta ggtaaatgaa tacatatttg atgcatggct | taaagattta ctttgacttt | tcaccccttt tacgtacgct aaatcaattc | 120 |
| gtaatttgaa tgaatcactt gcagcctgtt atttcagata | 19373 cgaaatgggt gccacacaat tcatcgtccc aacagttgct tcattggaat | aacaggactg acttcaaaag tttttgacct acttttaagc gaatggcacg | attatattta ggtaaatgaa tacatatttg atgcatggct attatgtcac | taaagattta ctttgacttt atggccaact | tcaccccttt tacgtacgct aaatcaattc atttacaagg | 120 180 240 |
| gtaatttgaa tgaatcactt gcagcctgtt atttcagata atatggcaaa | 19373 cgaaatgggt gccacacaat tcatcgtccc aacagttgct tcattggaat tgcaaaacaa | aacaggactg acttcaaaag tttttgacct acttttaagc gaatggcacg tggcactgtg | attatattta ggtaaatgaa tacatatttg atgcatggct attatgtcac tggccaaagg | taaagattta ctttgacttt atggccaact catttactgt | tcacccttt tacgtacgct aaatcaattc atttacaagg ggatgggaaa | 120 180 240 300 |
| gtaatttgaa tgaatcactt gcagcctgtt atttcagata atatggcaaa | 19373 cgaaatgggt gccacacaat tcatcgtccc aacagttgct tcattggaat tgcaaaacaa agtgactctt | aacaggactg acttcaaaag tttttgacct acttttaagc gaatggcacg tggcactgtg | attatattta ggtaaatgaa tacatatttg atgcatggct attatgtcac tggccaaagg | taaagattta ctttgacttt atggccaact catttactgt gaactcgttt | tcacccttt tacgtacgct aaatcaattc atttacaagg ggatgggaaa | 120 180 240 300 360 |

| <212> <213> | DNA Glycine max | • |
|-------------------------------------|---|-----|
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| tatctttaga | agctaggctt ctttgcatta gggatgtgca aaaaaatcgg gtcggatcga | 120 |
| accagattgc | aactgactcg actcgaacca gttgcgaaaa aaaacttgcc ccattttata | 180 |
| gtcagtttgg | ttcgacccga cccgttttgg caaaaaaaat tagtgacctg aacttgaact | 240 |
| gcattgctca | ctctcatcat cctacgatct ttntttctgt gcaacttcat ttgtttaaga | 300 |
| agggaattgt | ggtttcgaag ttgtacgacg aagcgattat ggtggatctg aaggaaacgc | 360 |
| tgagatacaa | tntctttctc ggaaccttcc ccggaatg | 398 |
| <210> <211> <212> <213> <223> <400> | 19375 429 DNA Glycine max unsure at all n locations 19375 | |
| | tntttctaag ttctttattt agtcttttta caanatactn gnccttcatt | 60 |
| | tgggcttggc ggccacgctc aacaaagtat tttcgacacc tactgtacgt | 120 |
| | aacgctgtta tgggaatgtt gcgacaatcc ttcaaaacct tattgataca | 180 |
| | ttggttgtca tgcggccata ccgaagtcct tctctatcat aagtcatcgt | 240 |
| | tttgaaatgt gatcaatcca tgttgctatg gctggactca gttcacgaaa | 300 |
| | ttttgataaa aaatgtgctt gcaaggagtg taggctgcat caaattagtt | 360 |
| | attttaagta tatattaaac ttaaataaac ttgaccatga tatatgaaat | 420 |
| cttacccaa | | 429 |
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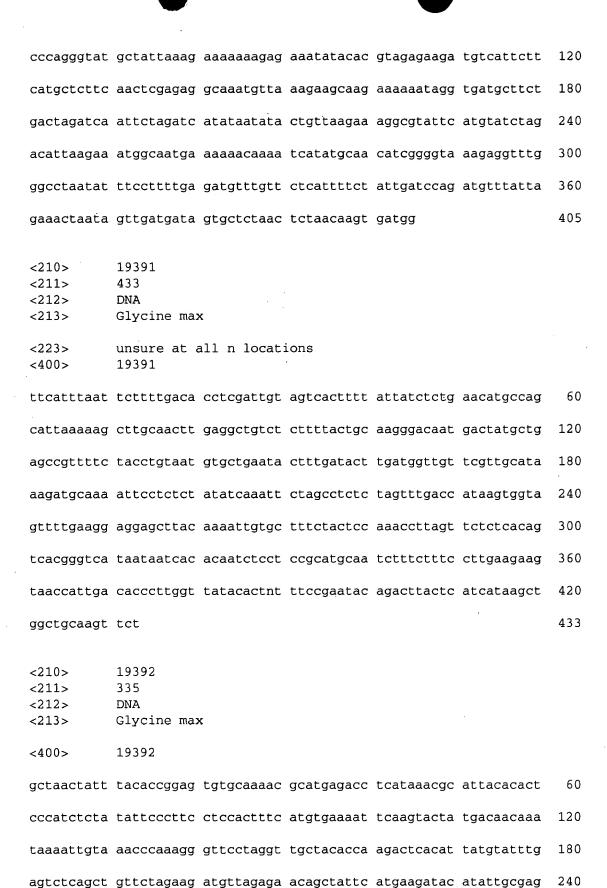
| gacggaaact | tragagteta | aaccttaaaa | gtaaatacac | ccaattntag | tgctccttta | 360 |
|----------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| gacggaaacc | ccagageeea | ggccccgggg | geadacacac | ccauçencag | egeteetea | 300 |
| agcctctgcc | gatcccacat | ggttggagca | ttntcacaaa | atagcgtgac | cctaactggt | 420 |
| ctccctatga | tttta | | | | | 435 |
| <210> <211> <212> <213> | 19379 398 DNA Glycine max | ĸ | , | | | |
| <223> <400> | unsure at a | all n locat: | ions | | | |
| agcttgtaat | caagtgaggt | ggagtggagt | tgataattca | gttagtgtag | actggttttt | 60 |
| ttctatttag | gagggatgct | acatgcttca | attcagactt | gatgtcgaag | cagagtaatt | 120 |
| ttggctcctt | taaattaaaa | ggggttcact | ttggaaggaa | aagtttaggt | tctcaaccat | 180 |
| ttgagtcaac | agttgatatt | atggtaaaat | atatgcacaa | ttggccaatt | gcacattcat | 240 |
| aacaaagtat | ggatttgtta | taacttataa | cttatccctc | actttagatg | ggtcgaatcc | 300 |
| aagaggaaca | tgcaatttaa | tcaattgtat | gttttgttta | agtcaatatg | cttaaagttt | 360 |
| tttcaacttt | tatccatgtg | cattangctc | ccctggac | | | 398 |
| <210> <211> <212> <213> | 19380 439 DNA Glycine max | ς | | | | |
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| tataacactg | caaaggtgat | gagaatagtc | ctcaaaattg | ttattataat | aacaaatttt | 60 |
| ccttttcttt | tattacatga | ccaatcaaac | tttagacaga | aaatgttact | aacactacta | 120 |
| gttaattaat | actatttaca | taatttcaaa | tcaggttaat | actcagacaa | ctataaaggt | 180 |
| caagaaggac | aatgaaggct | tagaatatgc | gttgaaatgg | caatgaaggg | aagagctgtt | 240 |
| tgtccaaatt | aaagatggac | taatcaccgg | catcattgac | ttgactgctc | gaaaactctt | 300 |
| ggatcaaaag | gttgtcacaa | atgattctag | tgattactta | tggtacatca | ccaggtaata | 360 |
| atattctatg | catgtccttt | ntaattaatt | caaagttgag | taatgttaac | ttctcctatt | 420 |
| aattatatot | tectacaaa | | | | | 130 |

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|----------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
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| acaatagcat | catttcttgc | actgaattgt | tgggagttgg | aagcaatctt | ctcaatcaga | 120 |
| ttcctagcct | caacaggagt | catatcacca | agagctccac | cattggcagc | atcaatcata | 180 |
| ctcctttcca | agttgctaag | tccctcatag | aaatattgca | gaaggagttg | ctcagaaatc | 240 |
| tggtggtgag | gacagcttgc | acacaatttc | ttgaatcttt | cccagtactc | atacaagctc | 300 |
| tctccactaa | gttgcctgat | gcctgaaatg | tcttttctga | tggcagtggt | cctagatgca | 360 |
| nggaagaatt | tctccaagaa | caccctctt | | | | 389 |
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| tataaatcta | agcttgcttc | tcaatctccc | cctttttgat | gatgacaatt | cttttatcaa | 60 |
| gaaagcatat | acaatttcta | ttgttcgttc | attcattcca | agcttcccct | ttgtcttgga | 120 |
| gattatgcct | aattttttg | aaatctaaaa | ctttatcttt | cttgatttct | ctaaacctcg | 180 |
| tttctctccc | cctttggaaa | catcaaaaag | ccaaagtgcc | caaaaaaaca | aatataattt | 240 |
| atccaggaag | agaacacaaa | accaatcata | ataccagagc | aatcaacatt | catacataat | 300 |
| tcaattatag | tgtatttaat | caaagaaaaa | tatccaaaca | aagaaaatca | atccaaaacc | 360 |
| ataaatatac | caagtcagag | gtcttataca | tagccaaaat | acacagcgta | gaaattataa | 420 |
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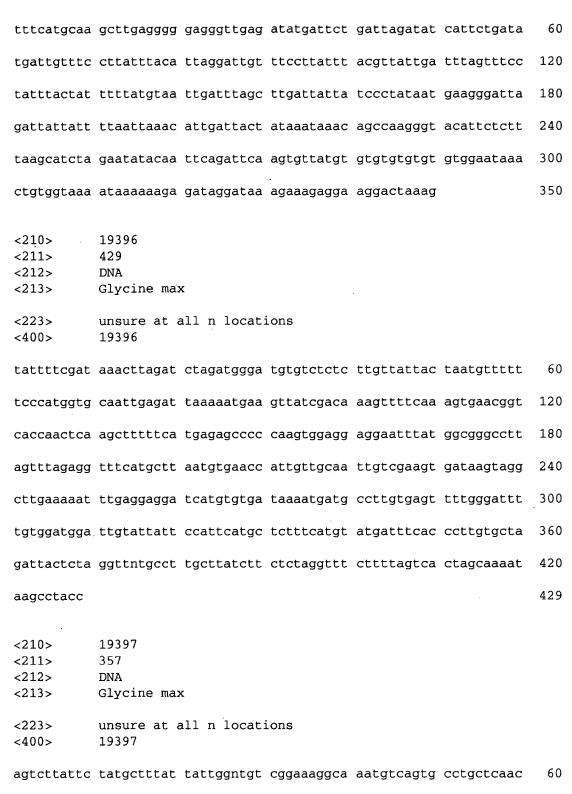
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| agagacttgc | aaggcaatcc ttctactgtt tcttggacag atactcaggt tacaatcaga | a 120 |
| ttacagtaga | teeteaggat caagaaaaa cagegtttae atgteetttt ggtgttttt | g 180 |
| cttatcaccg | catgttgttt ggtttatgta acgcccctgc tactttctaa agatgtatga | a 240 |
| tggcaattnt | tgatggcatg gtagagaaat gtatcgaagt ctttatggat gatttttcgg | 300 |
| tcttcggtgc | atcttttgga aattgcttag caaatttaga gaaagtgtta cagcgttgtg | g 360 |
| aagaatcta | | 369 |
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| gctacaattg | caccctcctt ctttcccttt tttgttaatt gtgtgaatga attagttaaa | 120 |
| ttgtcacagc | tttttgtgtt aattgttctt ttattgttca aagtgaagta atttttggta | 180 |
| acttttaatt | gtccgaaatc tcttaatctg aatatcttcc aagcaatttc tagttcttta | 240 |
| tagccaccag | aaagtgaaac gtcaacaaat gtaatataat aacaataaaa ttattccacc | 300 |
| attagttacc | attaccattt aacaataatg atagaaacat aaaaaattaa gactgattaa | 360 |
| tttttaattt | aaaggttaaa agtgaaaatt cataaaagtt taaagaccca aaacataatt | 420 |
| aatcctatct | ttttcatatg | 440 |
| <211> <212> | 19385 320 DNA Glycine max | |
| | unsure at all n locations 19385 | |
| cgagctcggt | gcccgctgat actctatagt gcaccagcgc gccttcanac ttgtatagtg | 60 |
| ngcataaaga | tatagatatg gggatagcat accttcaatg ctaaatagta cgatgaaatg | 120 |

| gtaccacgta | catttcctgt | aacagctgtt | aaagtattct | tgatactatg | tgatgagtat | 180 |
|-------------------------|------------------------------------|------------|------------|------------|------------|-----|
| agagataata | acgacgagct | attatcagcc | cattgcatga | gataatatat | aattatgcag | 240 |
| tatatttaat | atcacccggg | atatatagct | gattaatact | ttţactgcta | ccagagattt | 300 |
| ctttactaga | acgctattaa | | | | | 320 |
| <210> <211> <212> <213> | 19386 408 DNA Glycine max | ĸ | | | | |
| <400> | 19386 | | | | | |
| taatcaattg | aaaattgacg | gtgtgagatt | ctctctgttt | ctctactagg | ggcaattttc | 60 |
| cttggcaccc | tttatcatgt | tcaatttgtt | ggtaagttta | ggtcttttaa | tccaaaaaag | 120 |
| gaaacttggt | taccatgtga | gagtaatttg | gataaatgaa | aattgttttg | gttgttatat | 180 |
| gcatgagtat | ttcgatgctt | gtttgcaata | atgtaatata | caaaagtacc | taccacatag | 240 |
| agagtgccta | cgcaatttgg | aatcaagaag | tttcagattg | tgtgattgca | ttctctagca | 300 |
| ccaaagctat | tgcattgaaa | aattactgca | tacccaaaat | tactttaata | agttgcaacc | 360 |
| aatattactt | ggcaaaaaag | tagtctaaag | ctactctgtc | atcatgga | | 408 |
| <210> <211> <212> <213> | 19387 407 DNA Glycine max | c | | | | |
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| ccccttaggc | acttctctct | ctttcgaatt | tgcttggaaa | aattggttct | gtgaagaaaa | 120 |
| ttcaagccga | ggcgctctcg | aaacgtttcc | gtaacgtttc | cgtgagaaat | ttcatgaagg | 180 |
| tttcgaccgc | tcttcaagat | tcatcgctcg | gtctttgctt | tcttcagact | tcaacgggta | 240 |
| agtacctcaa | accaagcttt | tcaattcatt | ctatgtaccc | atggtggtcc | acatttcgtt | 300 |
| tcatgtatat | ttattcccct | tttcatttac | tttttatacc | cccttttgac | gtgcttaagc | 360 |
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| ttattcgttg | accacagagt | ggtacttgga | gatatgtcgc | gggggtcagg | agaccttggg | 300 |
| gacgtcaggt | ggggtgctat | tgcccaaaac | caagcttgac | caatcccgac | ccaacccggg | 360 |
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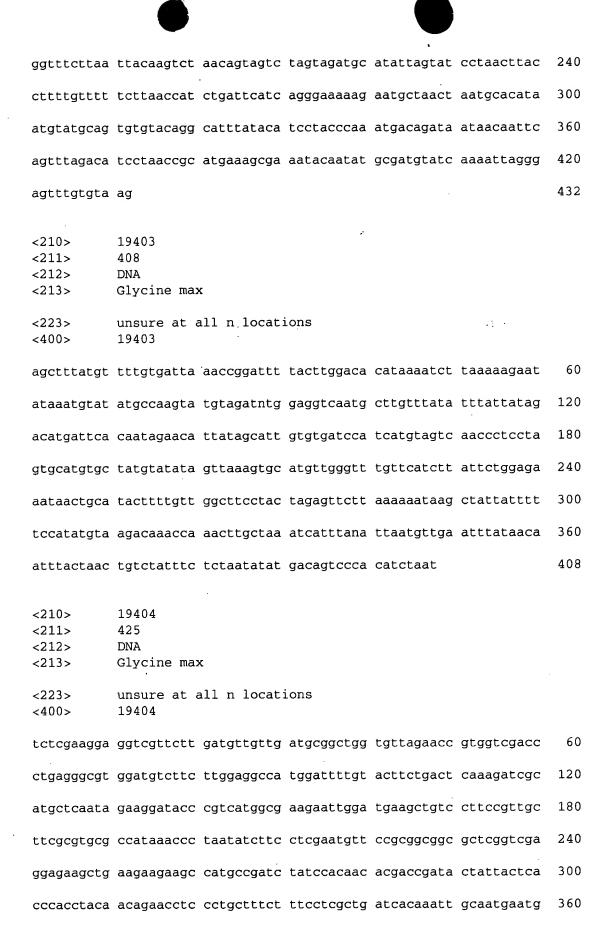
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| catttataat | agaccccctc | agcagcaaaa | ccaacctcag | cagaacaatt | atgatctttc | 240 |
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| tcggtgaaga | atttgtcgaa | tggactaatc | aattcacacc | attatccggt | gtgaatggaa | 300 |
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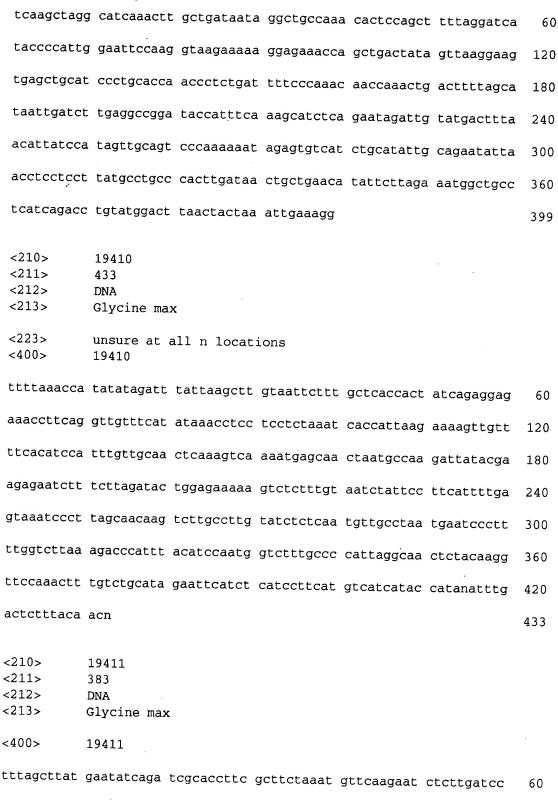
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| gaataatcta | a atctctttga agaagatgca tcagttcagt tgtcatgtgt aacaccattc | 360 |
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| tgatgtgcca | tcattttctt ctattttcta aacccttttt gctccattnt aattattgat | 120 |
| tggtcttaat | tgtcaattaa ttaggcagtt ttattatttg ggctcattta gctaatttga | 180 |
| tgtttctaat | ctaatttcaa gaattaatga aacattgggc ttaatccgga ttttggttgt | 240 |
| ggacttgaag | agggcaaata aagcagcgct taccttagtt aatttctaat taagaaattt | 300 |
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| caatatcaaa | acaaatctag aaacggagta acacaggaac tttggagtct tgtctcatat | 180 |



| tcgacggatt | cacctgtgta | atcctccgcg | ttactttctt | gttgctcttt | ttttctttnt | 420 |
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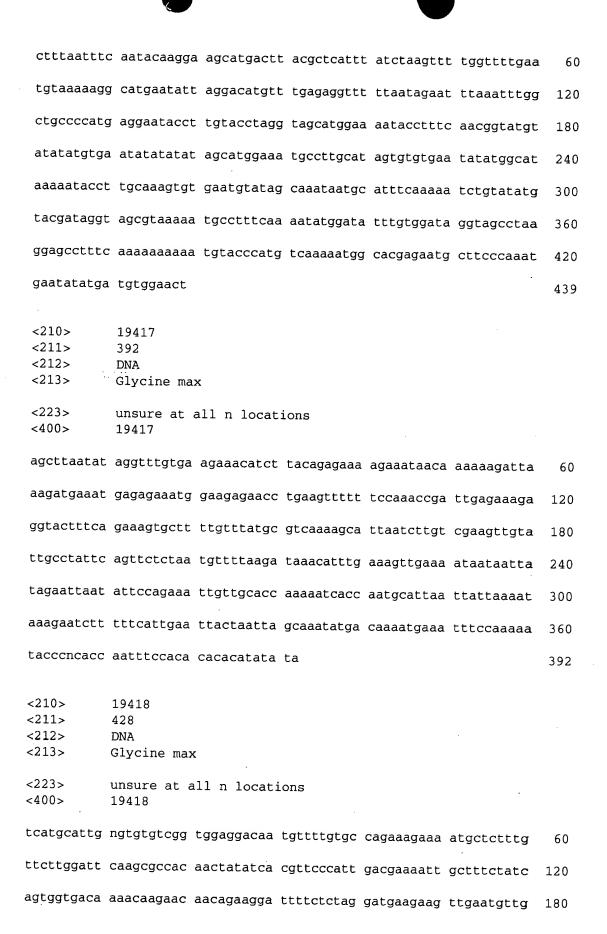
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| aaaagatgta | actcttcaaa agggttttga ctttttcaaa ttggttttaa gtttttctaa | 120 |
| aagttataac | tcttctaaat ggtcttcttg gccagacatg cagagtctat aaaagcaagg | 180 |
| ctttgatttg | cttttcaata cacttttcac attcattcaa tcaatccttt gcaagccttg | 240 |
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| tccaaacctt | gaaaacttgt gctattcatc ttttcattct cttctccctt tgccaaaaag | 360 |
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| gaacaaagaa | ct | 432 |
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| gctaagtgaa | cctctttgag aaaccaaacg tctctctggc tcacttagcg cggtggttcg | 180 |
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| acgttggata | gtctatgtga cactaaaatt gccaataatg cacctcacta acgcgttact | 180 |
| tttaaattta | acgacaagga ctattttgca aaacttatgc aaagataggg actatttttt | 240 |
| acatttcaaa | aagataggga ctaatttgta aaaagggtca taagtcaggg accaaaatgc | 300 |
| ttatntactc | gtacaataac acttgttcaa tgtttgacnt aaaaaattgt catgagacaa | 360 |
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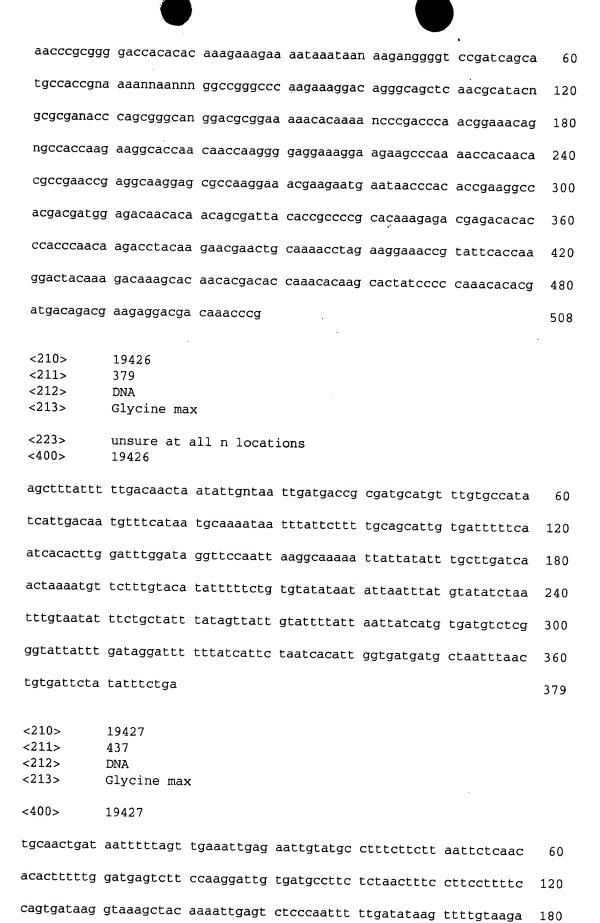


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| tgaaagaac | a ggaactgaat ttggaggaaa aaatagtttc tggaaaattc tccttatcca | a 360 |
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| gagaatgatt | t tcaagaatga gtcaacaagt tcaagatcaa gattaaagaa aagacatcaa | 120 |
| gaagaatcaa | a gattcaagaa taatcaagat caagatccaa gactcaagat tcaagaatca | 180 |
| agagaagaat | caatcaagat aagtatttaa aaagtttttc aaaacattga gtagcacaag | 240 |
| aagttttcac | aaaatcatta ccaaagagtt ttactctctg gtaatcgatt accagaatgt | 300 |
| agtaatcgat | taccagtgtt tntaaaacgt taagaatttt caaattcana atgaagagtc | 360 |
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| cattctccat | agcatcatca gtgtgttttg gttctgtctc ttagacgagt attgtctaac | 120 |
| cttgttgtct | gggggatgcc cttgtttgga ccttggttaa ggagtcacca atggttgagg | 180 |
| atccaatccc | atcatctact ctgagatccg taataaaatc attcaactct gatatttttg | 240 |
| tattaggctt | gatgtcatca aacttcacat tgatagcttc cttaataatt aaggttctag | 30Ö |
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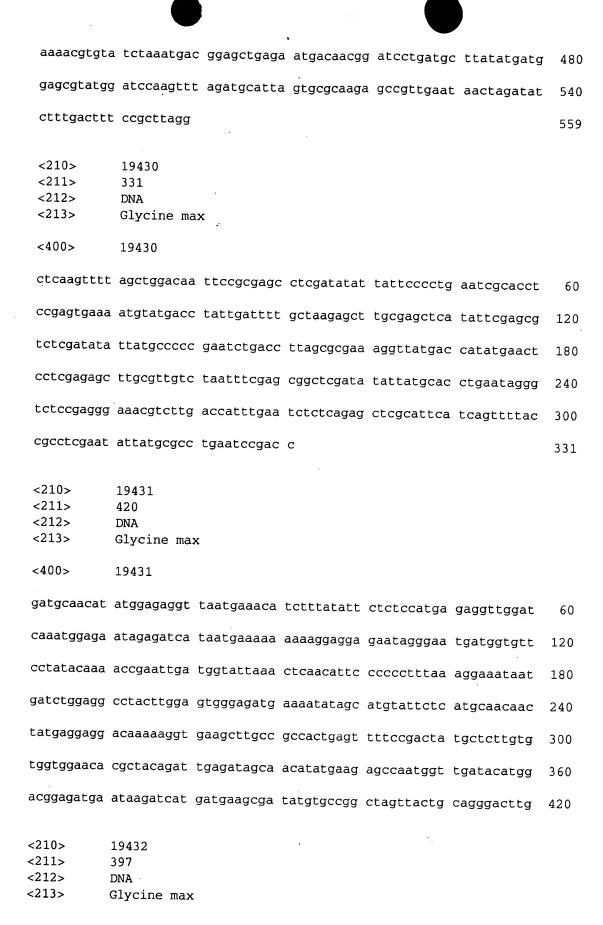
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| | gcgactggtc | c cctctcttcc cttcgcagct tgagttcact attgctaccc catagagct | c 120 |
| | cgcgaaattt | attccggcca tactcttcct tgcgagccct cttggtctct tgttcaagg | g 180 |
| | ctctggtggt | aattgcattc tcttcccgta atccggcata ctccttccgg atgtgtgta | g 240 |
| | cggccaactt | gaacttetet ttggcaagtt tegeetttee taactegett ttgagaget | 300 |
| 9 | ggacttcttc | gtcctcttcc ggtgcttcaa aactctcttc gctgacgact nttaacttg | g 360 |
| (| cgagccaatc | taaacctcgt. atatg | 385 |
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| | | ttcccttatt agagatgctc agacttagga tgtattcatt agagagctat | |
| | | ggcggtatta tcccggtgtg cgggttgtgt ctattactta tgcttgctac | 240 |
| | | ggactaaggt gtaacttact tatcacgctg acttcgcgct tagcgtgaac | 300 |
| | | agcacgcctt tgggcttctt tatgggcctt gttcaaccta agtgtgagtc | 360 |
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<210> 19423

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| ctacgcttc | gggttatatt gaaaataatg ttttcacggg acacaaaacc atatcatcct | 120 |
| gtacagtgta | a catgotaata ootactatgo aaaaaggttt togtaatata taottaaata | 180 |
| taacttttgg | tttattgtat ttcgtagttg atttgtttta atatattaag ttttaaaagt | 240 |
| ttgattttaa | a ttntttttta aatgtattat tttaattttt ttattaaaaa tattaatgtt | 300 |
| ttattaactt | ttaaatttta aaaaatatta aaacaatata tttcaaaaat ataaatgatc | 360 |
| ataataaaac | ttattatact aaaacaaaat aatcaggaaa cataatatat | 410 |
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| ttacaaggtt | tgcagcagtg ttgtgaaaat ctatctcatc gaggcattgg attagatgtc | 120 |
| aaatatgtgg | gtcatttatg ggtatcgcta atctatattt caatattaca taagtattta | 180 |
| ttattataat | ttattttgtt ggatattatc attgaatcat taagaaaatt cttgttatct | 240 |
| tttatgatta | agtacaaaaa ttatcaaatg gtcatcagta agtatgatga agatgtataa | 300 |
| aatacataaa | tattaaatca tatcaaaata ataactatta gttattatat atttttatca | 360 |
| agatcaaaat | aatattatat gagcatatta tacgctagta ttcaataatt ggatattaat | 420 |
| acgttaacat | att | 433 |
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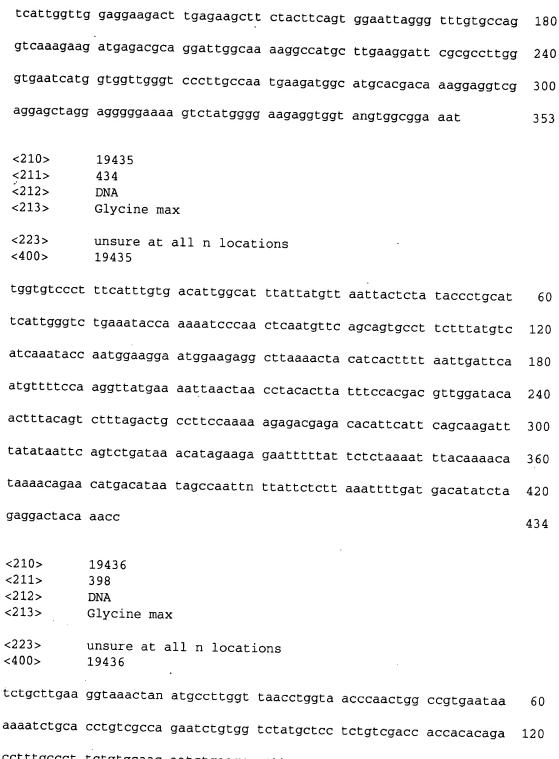
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| cttgaacgaa | a aatggttggc tcgctcctca ttgctctggg aatagataac gatctata | ata 360 |
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| attttttatt | catataaata tcaaaatatt tggaatttat tcctgtaaaa tttgattt | ta 120 |
| ttacgttagg | g catgitiget etetectaaa tataacaate tigataagge etataaat | at 180 |
| | tcaatttcat tctattagaa aaaggcccat gtgacaaaag aagtataa | |
| cgatcgaaat | gtttgaataa aatattacaa ttttttttcc atagatcaca tattcttc. | at 300 |
| taacactaag | aaaaaaaatc ata | 323 |
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| aagggtnatt | ttgatccctt gagaccgtgt anaccncaan ntaannctaa acggncnnt | g 120 |
| agccgtaaga | aatagactat aatctttatg tttcttatgc cttactgaca aaaggaatg | jc 180 |
| acgaaagagc | ttgatcccac ccatgctcaa cggcatatta tattagggag cacttttat | a 240 |
| atgaccgctg | actctggaca gtgcgataaa ctaccgataa ccatgacata gtgacgtat | g 300 |
| gcacgtgcat | atgacccacc ataactgtga agtcgaagat gaaaagaatt acggctgca | t 360 |
| gtgcctgtga | tgcttgtggc acaacctttg aattacaagg agacgactta tattacaaa | g 420 |



| <400> | 19432 | |
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| ttgttatca | t tgtctgaaaa taatgctcat caacataaca tttgtgtggt ttgcagaatt | 120 |
| | t agaagcatat tcccccaatg aatccgaaaa gtgttcaaaa cctattaaaa | 180 |
| | c tcactgaaaa atgatacaat aactgccaca aattttgtgt aggaaatcag | 240 |
| | tagagaaaaa acacatttaa tagcataaac aaacaattta taaagactaa | 300 |
| | c atgaacaaat catgagccag aaacccacgg atcaagaggg aataaaggat | 360 |
| catcaagaaa | a caatatgcga gtactttcat ttcaaga | 397 |
| <210> <211> <212> <213> | 19433 429 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19433 | |
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| | catgaatttg atgatgtcaa agaagaatct aacaaggctg cttcaaatga | 120 |
| | cttcaagaat aattcaagat tgcttcaaca aacaaagcct tgtttcaaga | 180 |
| | accaagcett geettaaaac aaagtgettt caagacatge aaggetetgg | 240 |
| | ccaggaagtg taatcgatta ccagaagaca gggttgagaa atagctgttg | 300 |
| | tgaatctgaa ttttcaacat gtaatcgatt accatatgtc tgtaatcgat | 360 |
| | gaaactttgg aaattcatat tcaaagtcat aaccctgcan attataactg | 420 |
| tgtaatcga | | 429 |
| | 19434 353 DNA Glycine max | |
| | unsure at all n locations 19434 | |

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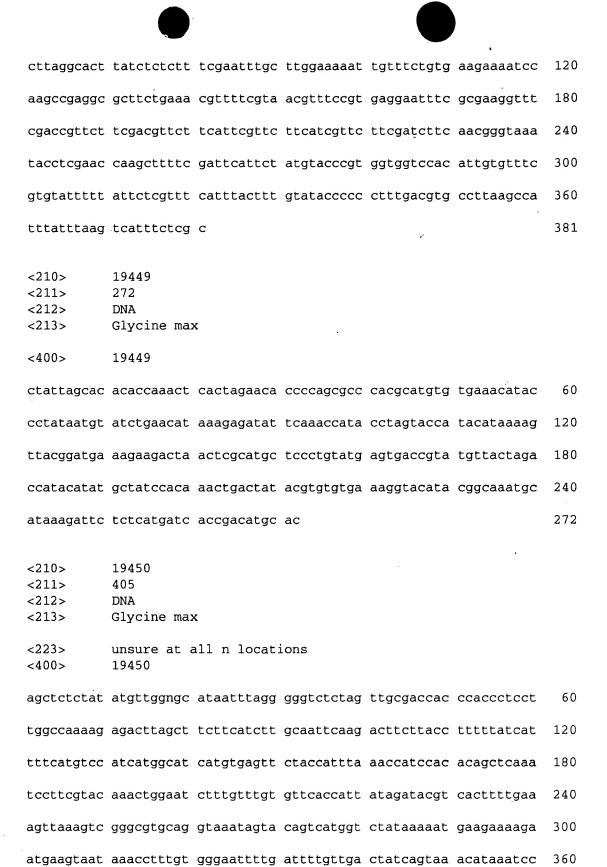
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| tnttgactg | a tacatngcan accngnaana anaaacncaa gcccccnanc nngggcaaaa | 120 |
| cgggacgag | g acgttagttc tatccnnggc aanncgacgc gcggaagcag cagaagccaa | 180 |
| cactccaaca | a gcaggtctgc ccagatacac gcaaatcaca gcaggggaga agcaccacac | 240 |
| ctgccccga | a caaacaccat tcgagactca acaccacacc gggaaaaatg ccagagggaa | 300 |
| tggagacaaa | a gaageceetg aegaagaeee cagaacaage agecaacaag etgacageae | 360 |
| acagcaggac | agcagectag atgaggaeca ggaacaecae geaaaecaag gegageggte | 420 |
| ccagaacgct | accgagggca ggccgcgaag tacacgaagc acgcctgaag caaagccaac | 480 |
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| | cacttttagt cttttctctc aaggtataca aggtgtttta agatcttagt | 180 |
| atcttttcaa | gaatttacag agatattttt acaagaaaga acgaaagaat gactcacgtg | 240 |
| aactgttcgt | gtattgtttc ttcaaagctt cttctatata tagtcttcgt ctccaagtat | 300 |
| ccattgtctc | tcaatgggtg gatccttcac tcta | 334 |
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| atagagaga | g agagagagaa | agagaaagtg | gcatggaaaa | ttgaaggaa | g aaagggagag | 120 |
| aagttgaac | t ttgaagcgtg | tctcacaaga | ctctcattta | tcaaagttg | t gacaagtgtt | 180 |
| acacatgtt | t ctatttataa | cttaggtcac | taactaaatg | aaattcact | t tttttgtgat | 240 |
| tttcatttt | atttcatgtg | aatctaagag | gaatattcca | aggatatac | c aaagaaatct | 300 |
| tagcatatto | c caagaatatg | ccaaaggctt | cttagcatat | tccttttaga | a tgccacaaga | 360 |
| atggaaggt | g tgactctagc | acatgggaaa | ggaatatgcc | tcaagaatat | gccaaaggca | 420 |
| tcttagcata | atccttgag | | | | | 439 |
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| attcccaatt | ttcaacttac | ctatttggaa | gtgacatcat | ggccgctatg | tcccagctat | 120 |
| ccattgtgga | ttcagtcaca a | aaacaaactt | gaatatgttg | gactatctaa | cacggngatt | 180 |
| ttcgattcta | tttccacaca (| gatgtgggaa | gcactttctc | aggttttgta | tttaaacctc | 240 |
| | atatccatgg t | | | | | 300 |
| | taagctcaaa t | | ggtaaattac | cctatctatc | aagtaatgtg | 360 |
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| gatacaccaa | atttgagaga t | tcccaatct g | aggaggaat (| cttcccgtgg | attccagtat | 120 |

| cagagaggt | c taggtgagtc aaggagctca | ttgcacaaag | g gaaagaagaa | attgccatac | 180 |
|------------|--------------------------|------------|--------------|------------|-----|
| cttctccaa | g taaatcattg tagctcaagt | caagatatco | g aagcttagag | agattcccga | 240 |
| tctgaggag | g aatcttcccc atgaatccag | taagagcgag | gtcgaggtga | gtcaaggaag | 300 |
| tcattgtcc | c atggaaagaa ggaattgaca | taccagetee | . aaataatata | ttgccgctca | 360 |
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| caagtctata | accttaatct aaacttgctc | aaactggttt | ttcacctaaa | attccaccaa | 180 |
| atcaaaattt | gactcctcaa cacctaattt | taccctagaa | atggcttttg | ccttcacttt | 240 |
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| caaactanga | ctaactcact ttaacctnca | atttctactg | aatccagaat | tagcctttca | 360 |
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| taccttaatg | caaaaaatgt catgccaatc | cctctaattt | agactaaact | cataagatcc | 120 |
| attcatgcac | acacgcatgt gtagaaaata | ccctattatt | tatatcaaca | tacaaggata | 180 |
| ttcaaaacat | tctagttacc atacatataa a | atttttttg | aaagaatact | tacacgcatg | 240 |
| ctcaaggtat | tgtgcccata tgttcatatc (| ctaaacattt | gctatttaca a | aactacctat | 300 |
| acatgtttga | aatgtatatc atacaaattt t | tattgtttc | tctcatattc a | atttatatgc | 360 |

| atgtcggaaa | gctaattaca | ttatgcacac | acttttgcat | ttaaaaggga | ctntcatgcc | 420 |
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| agcagaggca | aagagacgga | gaaagaagcg | agcgcgttga | ggaacacgcc | gttgagcttc | 120 |
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| aaaagcttac | tccgcacaat | ggtggcctct | tgagaatgaa | gcggttattc | ctccttttga | 300 |
| tgacgcatgg | acacttatca | ctgacctaac | tataattcgt | gcgacaggtc | ggccaatatc | 360 |
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| ccctatcgca | ccagatccaa at | ctagaacg | atgggtgatc | aagaggagac | gcatgaaçag | 180 |
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| agtatgaagc | agctcatgga ga | agaacgcg | gtcactgccg | ccgctgtcag | ttcggctgcc | 300 |
| gaaacaaaac | cgactctctt gg | caactacg | caccatcctc | ccttcaacat | agtacgacng | 360 |
| gtaagggaca | cactgat | | | | | 377 |
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| ataatgctca | agatcaggat gt | tcaaaatc | accaataaca | gaatgcacag | attcaccagt | 180 |
| aatggaatgc | tcagaatgat ca | aaaggtat | aaaatgatgc | ctaactaatc | tatgaaatgt | 240 |
| cctatctatc | tcaggatcaa ag | ggttgtaa | gtcagatgga | ttgcctctag | tcatacacta | 300 |
| cattcagcgt | gcacacaact ag | ttgccttg | tcatgtaaat | aaaggtgtag | gtttgaacta | 360 |
| cagctaccct | caaatgatat cc | aaatgact | tgaaatctta | tgagcaaccc | tataaaatta | 420 |
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405

ccatatatac ttgctaaaac aatggattag agctatttta gtcac

| | · | |
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| gtttgtttca | gacatgaaat gccattcata atttgtacac attaataaag aattgaagtt | 180 |
| ggaggtgatc | ctgatcgagc aaaacaacaa ataaatatgt cagatgaata tgagaggagt | 240 |
| gaacatgctg | atgatgactc ttcaatggga tcatcccaga aatgctcatc ctttgatttg | 300 |
| aatgaagaag | ctagtagcaa agataacaat gacaacgatg ataagggatt cgaggaagca | 360 |
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| gcatgagcgg | cgcttggcac ttgaacaaga tgtatatatt tatatatctc cctccaatat | 60 |
| gaattgtgac | tgtttaaatt tcatttgctg gtttgccctg tgctttgttg cttgttttca | 120 |
| atgtggcatg | tgatttaatt ttctgggcat tcttgggttt gaaattcctg acatcattat | 180 |
| tgctgttttt | ttgcattaaa ttgattgaac ccttgaattt aaattgtgta aaatattctt | 240 |
| gcaaaaacca | aaagttgttt ctaatgaatt ttccttgtgt attactacca tttatgagag | 300 |
| atgctggttg | ttaataatgt aacatcttta cattgattta tgatatggat aaaatgcatt | 360 |
| ctacactact | gatggaggtt tataaagaat tataattnga agtgttactt gctggaaggg | 420 |
| ttttccatat | | 430 |
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| <210> | 19453 | |
| <211> <212> | DNA | |
| <213> | Glycine max | |
| | | |

| <400> | 19453 | | | | | |
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| aattacgaat | taatagttca | aataataaaa | ttaaattaaa | ggaaattaat | atattaagat | 120 |
| tcaacgataa | atactttaa | tgcattttta | gtttaattat | ttattaacta | tttttaattg | 180 |
| aaaaaaatat | agtttgattt | aatatataca | tgttttgtgc | catgtaaata | ttaatattct | 240 |
| gtgatgtgta | tatttttcat | aaggtgtcat | aacatgttgc | ataggaatta | taacattgtg | 300 |
| attgagattg | gatgtatgtg | ataaatcgag | tatgtgttga | attgaagata | catgtgtata | 360 |
| agatettgae | gcattgagtt | gtgagctat | | | | 389 |
| <210> <211> <212> <213> | 19454 221 DNA Glycine ma | x | | | | |
| <400> | 19454 | • | | | | |
| acactctatc | ccctccactt | ctatgcaaca | cataacaagg | ctagttgtta | tagatactac | 60 |
| taataatgtt | gcatacagaa | actgaaccta | cggcttaata | acttatacgt | gccaagtcct | 120 |
| ttccatctgg | atagattatt | gataggggag | acaacaacaa | ccaatgccgc | acttgaacat | 180 |
| gtagaagcta | agactacgga | catgtcataa | acaatattct | С | | 221 |
| <210> <211> <212> <213> | 19455 432 DNA Glycine max | ĸ | | | | |
| <223> <400> | unsure at a | all n locati | lons | * 2 | | |
| gggataagga | cctactattc | catggactaa | ttatctttta | cttatttnca | tggttctgtt | 60 |
| aattttttt | ctacgaaaat | attctagcac | tcacgtatat | atggaccgag | aagaacatac | 120 |
| gaaaaatatt | atagcactca | tacatttaaa | gaagagtgaa | gaattgaaga | attcatgact | 180 |
| aaaataccat | gggtgatgaa | tctgagaagg | aagcattcat | ctctgagcgg | ccttaagaac | 240 |
| taattatctt | tgtaaactaa | tttagtagca | tatgtacata | ttatttgcac | tccaaataag | 300 |
| atattattt | ttttattgaa | 255222222 | 2021120111 | 2225455 | | 260 |

ggtccttaca ttaaaaaaaa taaaaattct cacatcattc tatgaatata tatatataa 420

| tatatatat | a ta | | | | | 432 |
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| <210> <211> <212> <213> | 19456 378 DNA Glycine max | × | | | | |
| <223> <400> | unsure at a | all n locat | ions | | | |
| ggcactcnat | cttacacaaa | ggntcatcct | tectetegeg | gaaactcagt | atatcagtat | 60 |
| tggcttcgct | tctttttaat | ctaggaagaa | acggggtaag | r tacttacttt | caacgtctcc | 120 |
| cttagtagta | aggctttgag | atggacgagt | attgagccac | : ttcaaggcat | ctccaattaa | 180 |
| agaataatgg | aaaactctaa | gataatggtt | ctccttttta | ttctgagcca | tccttattgt | 240 |
| actgcgtaga | tcatatataa | tgaccaggtg | gttgtatgga | tcttcatggc | caagtcagca | 300 |
| aatgcatgtt | gaccaaaaag | agggatggat | gtccgcttaa | cctctgttgc | tgggtattgt | 360 |
| ctcttggatt | gccatgct | | | | | 378 |
| <210> <211> <212> <213> | 19457 424 DNA Glycine max | | | | | |
| <400> | 19457 | | | | | ٠ |
| | gaaattgcac a | | | | | 60 |
| | ttgtacaagc 1 | | | | | 120 |
| | aacaaatata a | | | | | 180 |
| | tttttgcaaa a | | | | | 240 |
| | gccacatgtt a | | | | | 300 |
| cattaatttg | tggactattt t | ctcaaatgg | attgataaaa | agaaatagac | tattgaaatt | 360 |
| gctttctagg | tttgcaataa a | agcctttatt | ttgcaaagta | ctagtgttga | aaccttgctg | 420 |
| atat | | | | | | 424 |
| <210> <211> <212> | 19458 371 DNA | | | | | |

| <213> | Glycine max | | | | | |
|-------------------------|---|-------------|------------|------------|------------|-----|
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| tttgcttatc | gttcaactga g | gaagatatcc | ttttgttaag | ggtggattct | ttttgaggtt | 60 |
| agagtttttg | tttaagcatt t | ccttgcacc | cttcttccaa | aatttcaagt | gtcatggtta | 120 |
| aggattttac | cctcttttca a | aggtcaagac | attctaactt | gaattgccct | gatttcttgc | 180 |
| actcacaaca | aatgattgga c | ctgttttctt | tcttatgata | ggatcttttg | gacgatccgc | 240 |
| tgaacttgga | ggattccttg t | ttctccaca | tacttctgat | ctttcttgtg | ataagactta | 300 |
| tatcattatc | ttcttcatta g | gagtcatcat | gcttatattc | ttcttctaaa | gcatcattca | 360 |
| ccacaaàgac | t . | | | | | 371 |
| | 19459 417 DNA Glycine max unsure at al 19459 | l n locati | .ons | | | |
| ttgagaatcg | tcaaagacgg a | catgaaaaa | attttagtta | ctttacaata | ggatangcac | 60 |
| atagattggg | atttggaagt t | atgtcaacc | gcccactgct | ggacattgta | taacgcagca | 120 |
| tatctcaggg | cgctacttat g | gttacatat | tgagttgaac | agaggctcca | gaaatgaagt | 180 |
| ttaagcagtg | aaacgtaacc a | tcccgacta | gacagaacct | gagatacata | tttgttacgt | 240 |
| ctagacaaga | tacaagtccg a | acagcatat | attgtttcca | taaaattcaa | tatatggagc | 300 |
| agaacacaat | cagataactc a | ctcaacctg | tgcctatctt | ttatatttgt | catctgtgtc | 360 |
| ttcatcgctt | ttetttetet te | ccttcatca | tccattgata | ctgaatgtgc | tcggaca | 417 |
| <211> <212> <213> | 19460 386 DNA Glycine max | | | | | |
| ttatctttc | tgcttgagaa ca | attaggtct | ttgtacatgt | gcaactctaa | gcgacattat | 60 |
| gtaggaggta | gaaagattgt gt | ctaaaggt | acaaacttct | tcaatgctta | aaactctttg | 120 |
| ttataatcga | ttacaaggct ga | atcatattc q | gattacacaa | gtgtctgtag | cttgtagaga | 180 |

| gatțctagta | tcggattatt | catttaccag | ttaactgtaa | tcaattacgt | aattcatttg | 240 |
|-------------------------------------|---|------------------|------------|------------|------------|-----|
| agaccatgtc | tgagacttca | tgagtctctg | ctttcatcga | ttatcagata | atcgtaatcg | 300 |
| attactgaat | tcttaagatt | gttcccagat | gcgatctaga | acactttaat | caacttcatc | 360 |
| aataatctaa | tcgattacat | agttct | | | | 386 |
| <210> <211> <212> <213> <223> <400> | 19461 437 DNA Glycine ma: unsure at a | x all n locat | ions | | | |
| | | taggctcaaa | attttqataa | atattoottt | tcaaaatoot | 60 |
| | | | | | | |
| | | aaaaagcttt | | | | 120 |
| aggtttctgg | taatcgattg | cattattata | ttttgaaggg | tcatgacttt | tgaatttgaa | 180 |
| tttcaagagt | ttcattgctg | gtaataaatt | acagacatat | agtaatcaat | tacatgttca | 240 |
| aaattcaaat | tcaaaaccct | tttcaacagc | tatttctcaa | acttcccatc | tagtaatcga | 300 |
| ttacactgcc | tggtaatcga | ttaccagagt | cttggatgac | tttgaaaccc | tatgttttaa | 360 |
| ggcaaggctt | gatcttgaag | aaatcttgaa | gcacgactct | gtttgttgaa | gcaatcttgt | 420 |
| attaatcttg | aagcagt | | | | | 437 |
| <210> <211> <212> <213> | 19462 379 DNA Glycine max | « | · | | • | |
| <223> <400> | unsure at a | all n locati | ions | | | |
| agcttttcgt | acgatgtagc | actactgatc | ttaaaaatac | gtaaacggaa | ttaccgataa | 60 |
| cagacttcac | aaatttgtgt | cagtggtgac | tatcgagctc | ctcctcctgc | aattgccata | 120 |
| ttgatgctta | ttttgtctaa | cttcacctat | cgagaattgt | tgtgtcaagt | atccatttat | 180 |
| gttcaacctc | tcaaacatta | ataaatagaa | atctcattct | tacatgaata | catgcataag | 240 |
| taaccatggg | cttgattgaa | tatgaaggat | gtctcaaact | acacataggt | ntgtaaattg | 300 |
| ttttgtgctt | ttgcaatacg | cagtcactaa | actcctatag | ttaactagcg | gtatcaaaac | 360 |

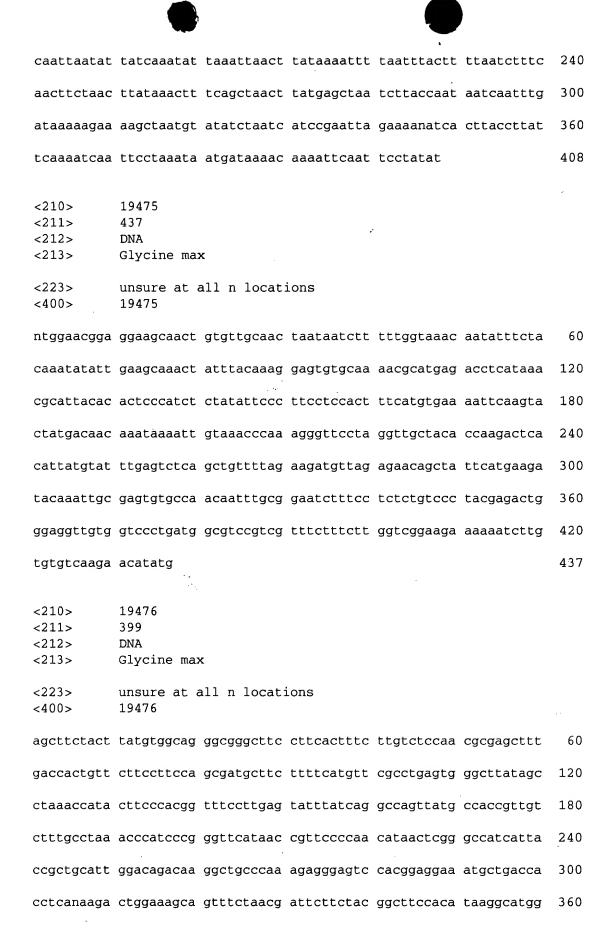
| atgttaatcc | : aacctaatt | 379 |
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| <210> <211> <212> <213> | 19463 426 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19463 | |
| tcccactgat | gagagtetge teacteatgt tgteaetttt etaatgattn ttteeeettt | 60 |
| tggattgaca | aatataggtt gttttttgtg tatatttcgt ctgattggtt tgtatcattg | 120 |
| atttctcttt | gcagataaag gatggatttg ctgagggcaa ggatcttgtt gtgtctgtca | 180 |
| tgtctgctat | gggtgaggaa cagatttgcg ccctgaagga tattgggcca aagaactagc | 240 |
| ttttggtgct | ggcagcctgt tgtttctatt taagcaaaga tccttttgta agcctttata | 300 |
| ttggtttgtt | caagacctgg cttatggctt atagattcta gtcagactag tcttaacaat | 360 |
| ggtgtttatg | gatgtggtca cagaaactat atcacatttt ttctggtttt ctatgctgtc | 420 |
| ctatga | | 426 |
| <210> <211> <212> <213> | 19464 327 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19464 | |
| tagcttctag | aaggagatca acttgatgtt ctatgcttct tgaaggtggc agtccatgag | 60 |
| gaatctcctt | ggaaaagaca tetttaaatt eetgeaataa gggttgaaca etaggagaaa | 120 |
| cataaatagt | taactgatta gaattatcac tetetetett ttgtgtatca etettteet | 180 |
| cgggtgtatc | actettett tteatattee tttgtggage eteactattt tettetett | 240 |
| gttctctctt | ttctctcatt ctgatttagg catcacatgc ttatctangg gatagagggc | 300 |
| taagaattaa | ccacgaagat ttgacta | 327 |
| <210> <211> <212> <213> | 19465 436 DNA Glycine max | |

| <400> | 19465 | | | | | |
|----------------------------------|---|--------------|-------------|------------|------------|-----|
| ctataaaact | cagcttacaa | ggctgcgagt | ggggttttt | ttatcttttc | ccttatgtta | 60 |
| tcaaacataa | aaagggaaaa | ggtaatattg | tagccgatgc | tctttctcgg | cgtcatgcat | 120 |
| tactttctat | gcttgaaaca | aaattgattg | gtcttgaatg | tttgaaaagc | atgtatgaaa | 180 |
| atgatgaaac | ttttggagaa | atttttaaaa | attgtgaaaa | attttcagaa | aatggtttct | 240 |
| ttagacatga | aggctttctt | ttcaaagaaa | acaaattgtg | tgtgcctaaa | tgttctacta | 300 |
| gaaatttgct | tgtttgtgaa | gcacatgaag | gaggtttaat | ggggcatttt | ggggtccaaa | 360 |
| agactctaga | aacattacaa | gaacattttt | attggcctca | tatgacatag | gatgtgcaga | 420 |
| aattttgtga | acattg | | | | | 436 |
| <210> <211> <212> <213> | 19466 405 DNA Glycine max | | | | | |
| <223> <400> | unsure at a 19466 | ıll n locati | ions | | | |
| agcttataat | ctntcttaag | tggggtgtat | tcaatcctga | attttaaaag | attttttatt | 60 |
| aaaaaaatta | caattttcaa | tcaaactttt | aaataataaa | aataagtctt | attggtacaa | 120 |
| atatttcaaa | ataaatatca | tgtgaagatt | taagacttat | ctgtttaagc | tttagagaat | 180 |
| aatggttcta | gtttctcttt | tttttttag | aagaaaccat | tttcacctca | taaaatatat | 240 |
| tgaaatagtt | ctttagttta , | ttttttctga | agacttgccc | aactattata | tttttgaaat | 300 |
| atatctttta | acttcaaaat | attttccttt | canaatcaaa' | taaacataac | cttanacaaa | 360 |
| aatattaaat | tntatttagc | agtttagaaa | gacatggttg | agttc | | 405 |
| <210> <211> <212> <213> | 19467 435 DNA Glycine max unsure at a | | ons | | | |
| <400> | 19467 | | | | | |
| | ggattataac | | | | | 60 |
| qaacaaaatt | cttttctaag | atgaaggaac | taaatttatc | tattttdaaa | agacggcatc | 120 |

| t | acactggtc | caagaggtac | cattttttta | agccattaga | tcaatctcgt | gcctacatta | 180 |
|----------------|--------------------------------------|------------------------------------|------------|------------|------------|------------|-----|
| t | tacaataca | ccatcaacac | cttacaattt | ctcaatcatc | ctctttcact | ttcctgtcca | 240 |
| С | actttcgct | aacgccgaca | gacacgtcgc | tgctgaacct | gcaacaggta | aacccttgcc | 300 |
| g | gcgacactt | tatcacacaa | gcaatttaga | atcatgattc | caacaatatc | acacacatga | 360 |
| a | acctaacac | ctcanaattc | accaccaatc | taaaccttgc | aattgcaaca | aagacacata | 420 |
| a | atcagttgc | atcaa | | | | | 435 |
| <; <; <; | 210> 211> 212> 212> 213> | 19468 216 DNA Glycine ma: | x | | | | |
| aç | gcttctatt | tatcagaatt | tattagatcc | tcattattta | ttagtttcta | aaacactaca | 60 |
| at | ctcctgac | ttatcaacca | aagaactttc | agctatctca | gaactttgcc | cateattact | 120 |
| tç | gaggtaatc | atactcagct | tatcagaact | tccctcttct | tttatgttgg | taactatttc | 180 |
| a | cctgattgt | agacactcta | agacacaatg | tatact | | | 216 |
| <2 <2 | 210> 211> 212> 213> | 19469 427 DNA Glycine max | c . | | | | |
| <4 | 100> | 19469 | | | | | |
| tç | ıtaggatta | tggagtaccc | atcacatgtg | gtacttttgg | gtttcgggcg | aaggtgcaca | 60 |
| ac | aagttgga | ccatccacaa | gacgcgcata | aacccaccat | cccctgatgc | ccacctccaa | 120 |
| ct | aagctcac | gtactaccac | gtagcccata | tggctgaact | ctctcaacac | caggtcccca | 180 |
| to | aatcctgg | aggacgggac | gacatcaaag | taattcagca | tgtaaacagc | acaagctatc | 240 |
| to | agccaaac | gaaacagggc | gaaggcagaa | aactctgccc | aaaacgccaa | ccagaatcaa | 300 |
| | | | | | tgcgattcgt | | 360 |
| at | ccactcga | aattatgact | gcaagtctct | agtacataag | cctacattca | gaccgtaggg | 420 |
| at | ctact | | | | | | 427 |

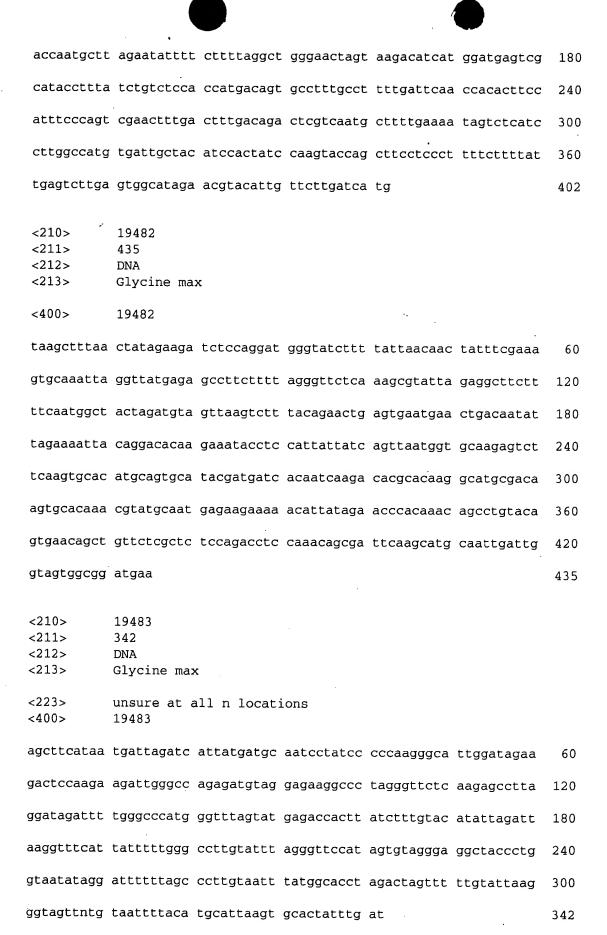
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| | Glycine max | |
| <223> <400> | unsure at all n locations 19470 | |
| agcttgagnt | tattatagnt gaccaataaa aaagctaatt aagtttttag tcatggggtt 6 | 0 |
| ttaaattttt | tatccctaaa ctattntaaa aaaaaaatta attcctaaaa aatttgttgt 12 | 0 |
| tattaaaagt | agttcatgtc attgatatcc tctattaaat aatgaaatgt tatgaaacaa 18 | 0 (|
| gttagatgtc | aggtgtatgt catatatatg tcacgtaagt tttattacat ttaagtaaat 24 | 0 |
| aattagaata | ttatgttatt aaaatgtaaa aaaaaaaaga aaaaagtcat tattgaaccc 30 | 0 |
| taacgcttca | agcaccatga actacattaa ccatccatga gcagttcatc ctcctcatct 36 | 0 |
| attatctatg | atgatattcg tgaccattta agtaggaatt atttat 40 | 16 |
| <210> <211> <212> <213> | 19471 437 DNA Glycine max | |
| <400> | 19471 | |
| tgaaccatga | atggaggcct ttatctatgt tttcatatat cttaaagggg tacaaaacaa | 60 |
| acctttcccc | tagtccctaa cccctctaag cttagaaaaa aattaggaac acgaaactat 12 | 20 |
| gctaagaccc | gctaatttct cgcttataat tcgctctata ttgagatcag aactggtgct 18 | 30 |
| aatctagaac | gaattcgaat ttatactttt gtgggtctct aaaatcaagt cctatctttc 24 | ŧΟ |
| aaatgccttt | ggtctcattc aatttagagt tccctacaat gttgtatgac tattatacca 30 | 0 (|
| aaacagattc | agagagacca atctgtagat gactttgcat gtagagggtt atttgagtaa 36 | 50 |
| gtacatggtt | tgtgatgcac aagatatatt ctcaagtatt tctctgtttg tcttggtaag 42 | 20 |
| gttcacgagt | gtccatt 43 | 37 |
| <210> <211> <212> <213> | 19472 385 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19472 | |

| agcttctagc | tttttcttgt | ctgccttaaa | ataaattgtt | tggctgctac | aggttcgcaa | 60 |
|-------------------------|------------------------------------|-------------|------------|------------|------------|-----|
| atggagagag | tggcaagctt | tgcaagacct | taaatcactt | ttttcagaag | aggcaagcaa | 120 |
| atgcttataa | agtttggtct | tttcttccat | gtttatcact | tctatttcag | tgtcacttat | 180 |
| tggttttctg | ataaatatta | tactctaggt | tcttgatgtt | attgtatcca | gcacaaatga | 240 |
| tttggggcct | ttcagtcttg | acaatttcag | aanaaatttg | tctgcttcat | ggagagttgg | 300 |
| attangaaat | tcaaacgctg | aacatgaggt | ttggaagaat | tatgtgcttc | tgcaattctg | 360 |
| catggaactt | ttaatttggt | aattt | | | | 385 |
| <210> <211> <212> <213> | 19473 437 DNA Glycine max | ς | | | | |
| <400> | 19473 | | - | | | |
| aacttgtagt | tgtatagcat | ttggttactc | atggctcaat | cttaaagtaa | tataataaga | 60 |
| atttaatgat | cgcatattaa | taaatgcaaa | aaaaaattta | catgatgaat | caattaattt | 120 |
| ttaattatta | gttataactt | ttagaatatt | tattataaaa | ggcaataaat | ttatgatgca | 180 |
| tgatccatgt | atgacttatt | ttctcaaatt | ataatacata | ctaggtttag | tcttctttt | 240 |
| ctttttttt | tttggtaact | aggtttagtc | atattttgct | catttaaatt | cagtgttgtt | 300 |
| ttagtttctt | caatttttta | ttttaattaa | tttttttctc | ttttatatta | aaaaattaat | 360 |
| tatttaattt | ttatttaaaa | ttcttaaact | accactttta | ttcagaatct | taaaaaactt | 420 |
| aagtaaatta | aattgtt | | | | | 437 |
| <210> <211> <212> <213> | 19474 408 DNA Glycine max | | | | | |
| <223> <400> | unsure at a 19474 | ll n locati | ons. | | | |
| agcttgaatc | tntatgttta | tgatcaatag | aattatgaat | atagacaatt | aattcactaa | 60 |
| atatatggtt | tcaattaatt | ctgaaaagtg | gatcagttcg | atttttcct | tttaagatat | 120 |
| aaatattata | aattttaaat | atatctttt | acgtcattta | acatttatca | attaacttat | 180 |



| aggatgggc | a gcttaccaag | g atgtcttcct | cgcctgaca | | | 399 |
|----------------|----------------------|--------------|------------|--------------|------------|-----|
| <210> <211> | 19477 432 | | • | | | |
| <212> | DNA | | | | | |
| <213> | Glycine ma | ıx | | | | |
| <223> | unsure at | all n locat | ions | | | |
| <400> | 19477 | | | | | |
| tgtagaatta | a tggggtaccc | atcacatgtg | gttctatgt | g geggteggge | gatggtgcac | 60 |
| aacaagttt | ccacatcccc | aaagcgcgca | taaacccac | atcccctgtt | gcccacctcc | 120 |
| aattgagctt | acgtactccc | acgtagccca | tatcctcgtt | tctctcaaca | ccgggtcccc | 180 |
| atcaatcct | ccaagcttcc | ccaacatcca | ggtaaaacaa | ı cattcaaaca | taacaaacta | 240 |
| tcacagcgaa | gaaaacaggg | caaaggcaga | agctctgccc | aaaacacaac | tcaaaatcac | 300 |
| agctttttct | cacttaaaga | ccccagtaac | atttccttcg | f ttccaattcg | ttaaccattg | 360 |
| gatcgactcg | aaaattttac | tggaagtctc | tagtacataa | gcctacattn | tgaccgttgg | 420 |
| gatctactag | aa | | | | | 432 |
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| <211> | 402 | | | | | |
| <213> | DNA | _ | | | | |
| | Glycine max | | | | | |
| <223> <400> | unsure at a 19478 | all n locati | ons | | | |
| agctttgttc | catataaaag | ttaattcaaa | caaaataaaa | ttgatatagt | tgtgacaact | 60 |
| tctagggtga | aaaagtttat | caaatgatct | atgagtgaag | caccattatt | aatccttaga | 120 |
| tttattatta | aaatatttcg | atggttcaaa | ttaaaattaa | ataattttaa | acaaacaaat | 180 |
| aaataaataa | ctgtatttct | atctttttca | ttctcacccc | acccccaaca | tctcaaagat | 240 |
| gttccacagg | aatctgggaa | atcttaaaaa | ataaaaattg | ttcatacatg | aaaaattgat | 300 |
| aaaacattaa | atacattntg | tccataaaga | aatgctctca | tgtgtgtgcc | tgtaaatcat | 360 |
| tttcatcacg | agaataacaa | aataatctaa | gtaactgaac | at | | 402 |
| <210> | 19479 | | | | | |
| <211> | 432 | | | | | |
| <212> | DNA | | | | | |

| <213> | Glycine max | |
|----------------|--|-------|
| <400> | 19479 | |
| taggcctaga | ggggatggac cttttcaggt tttggatatg atcaataaca atgcctacag | 60 |
| gttggacctc | ccagaagagt atggagtcag caccactttt aacatttcta atttaattcc | 120 |
| ttttgcaggt | ggaactaata ttgaggagga ggaactaaca gatttgaggt caaatcctct | 180 |
| tcaaggggga | ggggatgatg caatcctccc taggaaggga ccagtcacta gagccatgag | 240 |
| caagaggctc | caagaggatt gggctagagc tgttgaagaa ggccctaggg ttctcatgaa | 300 · |
| cctcagggta | gatttctgag cccataggcc aaggttgtgt ccaattatct ttgtacatat | 360 |
| tagattaaga | tgtcattata ttttgtcttt gtatttaggg ctccatgatg taggtagggt | 420 |
| accctagaaa | ta | 432 |
| | | |
| <210> | 19480 398 | |
| <211> <212> | DNA | |
| <212> | Glycine max | |
| <2137 | 017 01110 111111 | |
| <400> | 19480 | |
| tcttgcttta | a aaatttgaat taaaacgttt agaaattgtt ggtaatcgat taccatatat | 60 |
| gtgtaatcga | a ttacacagtg caaattttga attcaaattt taatagctgt tgtaaatcag | 120 |
| ttttggcca | tggtaatcga ttacatcctc tggtaatcga ttactagaga gtaaatctct | 180 |
| tgaaaaaga | c ttttttaact taaatttctt ggccaaacct tttgctactt caattggaat | 240 |
| tcccttcct | a tttaatgtaa tettettaag aetttagaga etgtettggt eateeatett | 300 |
| gaatatctt | t gatttetttg tettgaataa aaetttgaga aacatgtaat eetttggeaa | 360 |
| catcaaaac | a tcagcttgat cctttgtcta caacctac | 398 |
| | | |
| <210> | 19481 | |
| <211> | 402 | |
| <212> | DNA Cludino may | |
| <213> | Glycine max | |
| <400> | 19481 | |
| | a ttgctcttct tcatctttac ttgggctatc tcagaccttt tatgtttgtt | 60 |
| gtctaagat | t ttgcatacac ctccttcaaa gtgaagtgtg tagcttctct ccatcatttg | 120 |

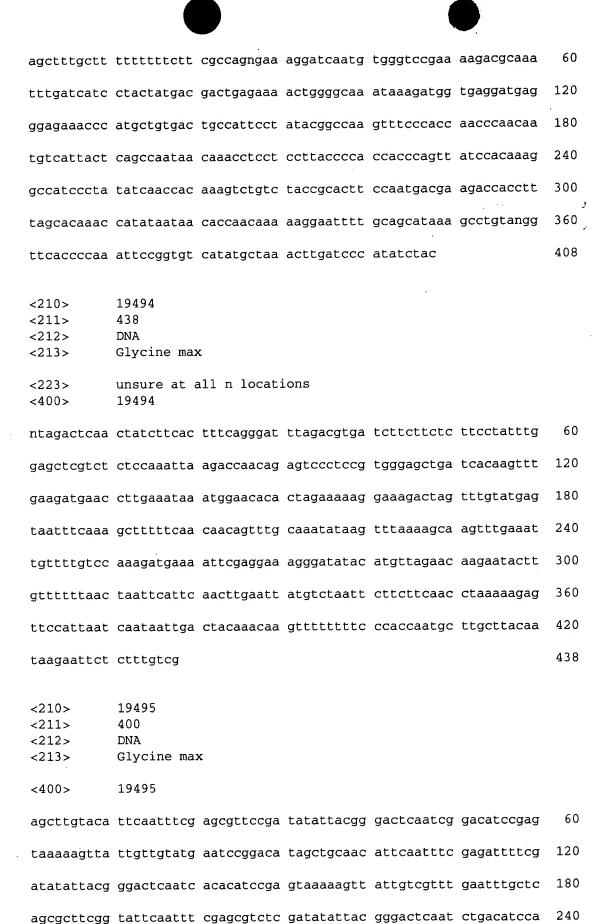


| <210> <211> <212> <213> | 19484 442 DNA Glycine max | |
|-------------------------------|--|-----|
| <223> <400> | unsure at all n locations 19484 | |
| tcatcaatgt | ttcttggttc aacctgtgaa acttaagctc atgttgttac acaaaattct | 60 |
| aagtctagag | ttcgtagaaa caccetttga gattteteea aetatgttgt etaaggggag | 120 |
| atccttttga | gttctccact ctctaggaaa ttctttaggc attgttgtat agatctcttt | 180 |
| gctttgttca | aaatcttcag ccttgatgtc atattcaaga aaaaaatcct tttcctaaaa | 240 |
| acctgtatct | tcttaaaaag aattttcttg aacaataaag ttagtttcat cacaaaccac | 300 |
| acatataaat | tcttcaacag ttaaagttct tttattgaat actccatatg ctttactatg | 360 |
| caatgaataa | ccaagtaaga tagcctcatc agcctttgca tcaaattntc caagagagtc | 420 |
| tttatcattg | tttaaaatga aa | 442 |
| <210> <211> <212> <213> <223> | 19485 383 DNA Glycine max unsure at all n locations | |
| <400> | 19485 | |
| | taaattagtc taaactttcg taagctattt aagctaagtc tagtccaaca | 60 |
| | gaggacgaag cttagtttaa gttagtctaa acctatgagg gctgtctaaa | 120 |
| ttgagcctag | tccaacaaga gggatctgaa gacgaagctt ccattcattc aatctcacta | 180 |
| gggatcgagg | tttagtaatt tatgcttcag catacaacac aaaagcatga ttgattagag | 240 |
| aaacatcttt | atatacatca gctggtttgt tagaaagacc caacatcttt acctattgct | 300 |
| tgtcaattta | cttacttgca ttngtactgt ttttagccta aacttagtta attctgtcta | 360 |
| aatcatcaat | catcaatgtt ttt | 383 |
| <212> | 19486 201 DNA | |

| <223> <400> | unsure at 19486 | all n locat | ions | | | |
|----------------------|------------------------------------|---------------|------------|--------------|--------------|-----|
| actatagaaa | ctaagcttgc | acttgagggn | gacatacaa | g ttttggttgg | , ttttaaacac | 60 |
| acccacgggg | ggagcttcta | atgaaattga | gccaacacca | a ttttcatatt | gttaatttaa | 120 |
| ttacctttgc | atgaggaact | aatattgcgg | tggaggaact | atcatcattg | tggtccaaga | 180 |
| ctcttttagg | gggagggaa | g | | | | 201 |
| <210><211><212><213> | 19487 397 DNA Glycine max | · x | | | | |
| <223> <400> | unsure at a | all n locat | ions | | · | |
| agcttttgat | aaagaaagaa | gaagaagaag | ttcaaagaga | ctcagaaatc | aatgtggaaa | 60 |
| aattgcttgt | gtaaagaatg | aattagaaaa | gatagatctt | aaaatgcaaa | acaaagcctt | 120 |
| gcttttatag | actcttcatg | tctggtcaag | agaaccatta | gaagagttat | gacctttaga | 180 |
| aaaacttaaa | accaatttga | aaaagtcaaa | aactatttga | agagttacat | cttttgattt | 240 |
| gttcagaaac | tatcactggt | aatcgattac | caaatcagtg | taatcgatta | cacaaagctt | 300 |
| ttttgtgaaa | ggatgtgact | cttcacaatt | taatttgaat | tccaacattc | aaacacactg | 360 |
| gtaatcgatt | accanatcat | tgtaatcgat | tacaaca | | | 397 |
| <211> <212> | 19488 430 DNA Glycine max | | | | | |
| <400> | 19488 | | | | | |
| tatccttttc | ttttacagat | tgcaacttta | aatacctttc | tatcttcatt | aactagccca | 60 |
| tcagatttcc | tcccttgaat a | attggcattt | caagagttct | ccgccaattt | cttctttgtg | 120 |
| aagcattccc | cttattctct (| cctattgaat | tgtctcgcaa | cctgcccttt | ggcgggggtg | 180 |
| tgaaaaggcc | tatcgaatgg (| gccaagggtg | catcttccgt | tgaaggaaaa | tgtgtggagt | 240 |
| cgccatcaac g | ytttatttga (| ggaaaacgtc a | agaaaaacca | aaatggaaaa | ggccgagggt | 300 |
| ttgcgtgttt t | gaaaatgag (| gattcgaaag (| ttgtttacgc | aagaggaagg | tattaacacc | 360 |

| cccacacac | c cgtcacaagg | gaaggcagco | : tctaataga | g tgtgaaaati | atgacttcaa | 420 |
|-------------------------|------------------------------------|-------------|-------------|--------------|------------|-----|
| aactattta | t | | | | | 430 |
| <210><211><212><213> | 19489 405 DNA Glycine max | ζ | | | | |
| <223> <400> | unsure at a 19489 | all n locat | ions | | · | |
| agctttatt | a gagagacaag | cctgttggat | tagagctaga | ı ttangtgcta | cgcactcttg | 60 |
| atttggatg | g gattgttaag | ctcttcaatt | tggaggaagg | ggttgtcttt | actgtcaata | 120 |
| gaactccca | a tatcttgctc | tattggatag | tatttactct | acaatgaccc | tgataatgat | 180 |
| tatgatttgg | g cttatgtgat | ctctagatgg | ttggatatgt | agaaatccta | ttagtactag | 240 |
| tateetetea | a aaataagaaa | ctagacgatg | tcattattgg | taccatacca | aagcatattg | 300 |
| acaatgtata | acccctagat | acctatactt | cataaggtcg | agtattgcat | atttgtctta | 360 |
| gatttttaga | atgtgagttt a | aagcctaact | caaccccaaa | gctag | | 405 |
| <210> <211> <212> <213> | 19490 442 DNA Glycine max | | | | | |
| <400> | 19490 | | | | | |
| ctaagcttcc | gtaaagaagc a | agaaacaata | gtggctcgtt | gttttctgta | tgatcagtta | 60 |
| cttcactttc | ttatcttcta g | gattettett | cctatgcatg | tctttgatct | gcataatctt | 120 |
| caactgttta | taatttttta a | tcttaaata | ttttaatgga | caacattata | caaaaactta | 180 |
| ataaaacaac | caccaaattt t | ttgataaaa | aatgcctttt | ttccctctct | catttactta | 240 |
| catattttc | tctctcttt t | attttgatt a | aatgtatacg | tggatcaata | catgattttt | 300 |
| ctctctcaac | taggttgggg g | tcacacctt a | acaacatctc | cttcctgatt | tgattttctc | 360 |
| tctcagctgg | gttgggggtc a | cactttaca (| ccatcttctt | cctgatttga | ttgatgtata | 420 |
| cgtggattaa | tacaatgcat a | t | | | | 442 |
| <210> | 19491 | | | | | |

| <211> <212> <213> | 395 DNA Glycine max | |
|----------------------------------|--|-----|
| <400> | 19491 | |
| agcttgagtt | gttaggtgaa gatacataaa tgatttcatt cattctatct tcttattttc | 60 |
| ttttcttgta | agtacttatt gagaggttta ttcaaaaata tcatagcagg caacattcct | 120 |
| agactcatcg | taacagatgc agcaaaacat ccatgccatg attaatcaag aatttaaaat | 180 |
| aaaaaaaat | aggtaagtat aatcaacact tcatagtata aaccttacca ccaaattagg | 240 |
| aaacaagttt | aatgagtcaa tagaggccag atcattaatg ttgttattag ctgtatgatg | 300 |
| aattgaaaga | cattagctgg tatccacaat ccagatagca ttgcatatgt tgaaaatgat | 360 |
| acttatcaag | aaggaaagta taatattacc caata | 395 |
| <210> <211> <212> <213> | 19492 437 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19492 | |
| ntaattcatt | atctttgaag acactgactt gcttccattn ttttaatcta gaactggaac | 60 |
| aagtggagca | agacccagat ttcataatgt tgtctcgcca atttaaagct actgcaatgg | 120 |
| agcttatttc | tgttttggag ctggtaaagc tttgaccttt aacacttcta ctaacattga | 180 |
| ctgaaaaaca | tgacatgcta tttgagcatt gagctttaaa aactctagta ttcttttaag | 240 |
| attggttgga | taatatcatt atactccatg tttgaataaa atattttgtg ttatagatat | 300 |
| gaactttttg | ttttatagtg ataaactata gtttttaaga ttttgggaat gccatccttt | 360 |
| ctgtgttgca | tttgtattag ttgcttacat ttccttgctt tattatagaa acaaaaaact | 420 |
| ggtggtagga | caaaagt | 437 |
| <210> <211> <212> <213> | 19493 408 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19493 | |



<210>

| agtaaaaagt | tattgacgtt | tgaatttgct | caaagcttcg | gtattcaatt | tcgagcattt | 300 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| cggtatatta | cgggactcag | tcgaacatac | gagtaaaaac | ttattgtcgt | ttgaatttgc | 360 |
| tcagagcttc | aacattcaat | ttcgagcgtt | ttgatatatt | | | 400 |
| <210> <211> <212> <213> | 19496 449 DNA Glycine max | κ | | | , | |
| <223> <400> | unsure at a 19496 | all n locat: | ions | | | |
| ctcagcttct | atataagctg | aaccattnta | tcaataaaga | caagttgagt | tttattcaga | 60 |
| aaattagagt | ttatctcttt | tatcttagtg | agagtgattc | tcctaaattc | ttgagtgatt | 120 |
| caagaacacc | ttggctgtat | caaaggactt | tcacaacctt | tgtgtgttgc | cctcgctgga | 180 |
| aagagtgatt | • | tttcatcttc | acccttgttc | tttcaaacca | caattccaga | 240 |
| aaatccacct | ctgcccagaa | ttatctcgtg | gccataactc | ccattttacg | cactcaaatt | 300 |
| aagtgattct | tgagcctaaa | ttgaattcca | aaacgagagc | ttccacctcg | ttttggaatc | 360 |
| acctcatttg | gagccctgta | gcttccgtta | ttgccatttc | tatatttctg | tccagccacc | 420 |
| acttaaccta | cgttntacca | tcccattca | | | | 449 |
| <210> <211> <212> <213> | 19497 316 DNA Glycine max | × | .: | | | |
| <400> | 19497 | | | | | |
| tcaatcttgc | tctaaattca | catggatgtg | agtatttatg | ggaggaggtt | gtatgtcatt | 60 |
| tctgttttaa | gagtagtgtc | ccgctggtaa | aactaacttt | ccaaatgttt | gccttcgcag | 120 |
| gaaatggccc | cgaggaagct | tgcctcaaag | aggtccagga | aggacaatgc | agcagaagga | 180 |
| actagttccg | ctccggagta | tgatagtcac | cgctttatga | gcgcggtaca | ccagcagcgc | 240 |
| ttctaagcca | tcaaggtgtg | gtcgtttctc | caggagcgac | gcgtccagct | caaggacgac | 300 |
| gagtatactg | atttcc | | | | | 316 |
| | | | | | | |

| <211> <212> <213> | 431 DNA Glycine max | |
|----------------------------------|--|-----|
| <223> <400> | unsure at all n locations 19498 | |
| tctcccctat | t tntgctataa atagggggag atttgattat gaaaggggtt cagcccctta | 60 |
| ggcacttctc | c tetetetete tegaaattge tgaggaaaat tattteegtg aacaaaatet | 120 |
| aagccgaggc | gctgccgcaa cgtttccgta atgtttctgt gagtaattac gtgaagattc | 180 |
| tcgaccgttc | ttcaagattc atcgttcgtt cttcgttttc ttcagtcttc aacgggtaag | 240 |
| tacctcaaac | c caagetttte aatteattet atgtaceegt ggtggteeac attttgttte | 300 |
| gtgtatttt | attctcgttt tcatttgctt tttatacccc cttttgatgt gcttaagcca | 360 |
| tttatttaag | f tcatttctcg cttaatctaa aaataaaata aatttccacc gatcatttaa | 420 |
| attgtatcat | : c | 431 |
| <210> <211> <212> <213> | 19499 398 DNA Glycine max | |
| <400> | 19499 | |
| ttatatttgt | gggcggacga accctgatga ccacagtggc atgatcgctg gctgacctac | 60 |
| accgatgttg | tcagcagcct cttcacgggt cctcacactg accettecte ctactgaage | 120 |
| atgtcgcctg | tgctaggcta gtgggcgcat accatctata aacatacacc aatgacttgg | 180 |
| ctgacaccac | ccacgagcgg gggtctttct cgcctatcct ttagggctcg acgacactcc | 240 |
| actgagggac | tggtctcggt actgatgata tgactatagt gctactttac cttacgcagg | 300 |
| atatgactgt | aagacatgtt gctgctaaca ctgttgatca acttggtacc cagagtacgc | 360 |
| actcgtacgc | ttaaatgagt ggagacacct ttgggctt | 398 |
| <213> | 19500 410 DNA Glycine max | |
| | 19500 | |
| cgcccccggt | gacgaagaca attgaataag cctctttttg cttttcacag gcgtcaacga | 60 |

| ttctcaaaa | t gaagggtctg | tgtttggaag | g tggcgatga | g gtggagaaca | acctcggtgt | 120 |
|-------------------------------|---|---------------|--------------|--------------|------------|-----------|
| cggaggtag | t gttgaagatg | gacccgctgt | cctcgaggt | t ggttcggagg | gtgcggtagt | 180 |
| tgacgaggt | t gccgttgtgg | gccacgccga | a cggagccgaa | a gcggtagccg | gcaacgaagg | 240 |
| gttgcacgt | tttgagcatg | gattggccgg | g cggtggagta | a teggaegtgg | ccgatggcga | 300 |
| ggctgccgg | g gagctggtcc | agcttcgact | ggttgaacad | c gtcggaaacg | agaccaacgc | 360 |
| cggtgatgga | a ttggaggacg | ttgttgtgaa | ccgtaacgat | tccggcgcct | | 410 |
| <210> <211> <212> <213> <400> | 19501 364 DNA Glycine max | | | | | |
| agcttctcaa | ggaagcttct | taaggaagtt | ttctcaagaa | agcttctcaa | ggaagctacc | 60 |
| tagtctataa | ataaaagcat | gtgtaacact | tgttgtaact | ttgatgaatg | agagtcttgt | 120 |
| gagacacaaa | tcaaagttca | acttctctct | ctttttctt | ccttcaattt | tgtgctcccc | 180 |
| cctctttctt | ttcctccatt | gaaacatcct | tccaagcttc | ttatccaagg | ctcatcttgg | 240 |
| tggtgaagct | ccttcttcca | tggcttattc | cctagtggat | ggcgcctcct | ctcacctctt | 300 |
| ctcctttgtt | ttccgctgca | tctccatggt | gaaaaattac | cattaaagga | cctcattgaa | 360 |
| gctc | | | | | | 364 |
| <210> <211> <212> <213> <400> | 19502 432 DNA Glycine max 19502 | | | | | |
| | gtacttaacc a | aggtecatet | tottatattt | accadatadt | ataastaaaa | 60 |
| | ttagacggtg (| | | | | 60 120 |
| | tttcataggc o | | | | | 180 |
| | cgtcatgttg c | | 4.3 | | | 240 |
| | gaggccttcc t | | | | | |
| | ttccaaacgc c | | | | | 300 |
| 3 3 | | o o o gu cu a | | aacggacgga | ciggiltttg | 360 |

| cgtaagagtt | ggaataacgg | ctcacaatta | gçggtgagtt | gtgatatgaa | tctggcaata | 420 |
|---|--|---|--|---|--|---------------------------------|
| taattcaaac | gt | | | | | 432 |
| <210><211><211><212><213> | 19503 399 DNA Glycine max | ζ | | | | |
| <400> | 19503 | • | | | | |
| agcttgctct. | tctgggcgag | ctaagtggca | agctcctccc | ctattttgcc | ataaataggg | 60 |
| ggaggagtga | agaagaaaag | ggttcagcct | ttttggcact | tctctctc | tcgaaattgc | 120 |
| tgaggaaaat | tatttctgtg | aagaaaatcc | aagccgaggc | gcttccgtaa | cgtttccgtg | 180 |
| agtaattacg | tgaagattct | cgaccgttct | tcaacattca | tcgttcgttc | ttcgttttct | 240 |
| tcagtcttca | acgggtaagt | acctcaaacc | gagcttttca | attcattcta | tgtactcgtg | 300 |
| gtggtccaca | tcttgtttca | tgtattctta | ttctctttat | catttgcttg | ttataccccc | 360 |
| ttttgacgtg | cttaaaccca | ttatttaagt | cacttctcg | | | 399 |
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| <210> <211> <212> <213> | 19504 431 DNA Glycine max | ĸ | | | ٠. | |
| <211> <212> <213> <400> | 431 DNA Glycine max 19504 | | tattttette | tattcaacaα | ccttctggag | 60 |
| <211> <212> <213> <400> tgccacccag | 431 DNA Glycine max 19504 ctcgcccagg | cgagcaaggt | | tattcaacag caccccatt | ccttctggag | 60 |
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| <211> <212> <213> <400> tgccacccag gaatcttctg acacccctt | 431 DNA Glycine max 19504 ctcgcccagg gagggcccaa gcctttttt | cgagcaaggt gtgggtctgg gggtgattct | ttgctatttg | cacccccatt | tttactaagt | 120 |
| <211> <212> <213> <400> tgccacccag gaatcttctg acacccctt ttcgtaacga | 431 DNA Glycine max 19504 ctcgcccagg gagggcccaa gcctttttt tacttgttt | cgagcaaggt gtgggtctgg gggtgattct ctttccgtaa | ttgctatttg tttttcgtaa cgttacggaa | cacccccatt | tttactaagt acttacgaat tacataatca | 120 180 |
| <211> <212> <213> <400> tgccacccag gaatcttctg acacccctt ttcgtaacga tcccttttt | 431 DNA Glycine max 19504 ctcgcccagg gagggcccaa gcctttttt tacttgtttt gacttacgga | cgagcaaggt gtgggtctgg gggtgattct ctttccgtaa atgttacgga | ttgctatttg tttttcgtaa cgttacggaa acctcactaa | cacccccatt agttacggaa ccttgtggat | tttactaagt acttacgaat tacataatca atgcttccat | 120 180 240 |
| <211> <212> <213> <400> tgccacccag gaatcttctg acacccctt ttcgtaacga tcccttttt ttgatttccg | 431 DNA Glycine max 19504 ctcgcccagg gagggcccaa gcctttttt tacttgtttt gacttacgga gtatgtcatg | cgagcaaggt gtgggtctgg gggtgattct ctttccgtaa atgttacgga gaaccttacg | ttgctatttg tttttcgtaa cgttacggaa acctcactaa gattgtgcat | caccccatt agttacggaa ccttgtggat ttgtgcaacg | tttactaagt acttacgaat tacataatca atgcttccat ttttgttttc | 120 180 240 300 |
| <211> <212> <213> <400> tgccacccag gaatcttctg acacccctt ttcgtaacga tcccttttt ttgatttccg | 431 DNA Glycine max 19504 ctcgcccagg gagggcccaa gcctttttt tacttgttt tacttgttt gacttacgga gtatgtcatg cggaatttca | cgagcaaggt gtgggtctgg gggtgattct ctttccgtaa atgttacgga gaaccttacg | ttgctatttg tttttcgtaa cgttacggaa acctcactaa gattgtgcat | caccccatt agttacggaa ccttgtggat ttgtgcaacg caatattttc | tttactaagt acttacgaat tacataatca atgcttccat ttttgttttc | 120 180 240 300 360 |

| <212> <213> | DNA Glycine ma | x . | | | | |
|----------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| <400> | 19505 | | | | | |
| tctagcttct | ccctcttttc | cctataaata | gggggaggag | ggaagaacaa | aaacgttcaa | 60 |
| ccctcctggt | atctgaggat | cacttgaaat | tagtgaaaaa | aatcgtttcc | gagaagaaaa | 120 |
| tcgaagccga | ggcgcttccg | taacgcgtac | gagacgtttc | cgtggttgat | attgtgaaga | 180 |
| acttacgcca | tccttcgttt | gctatacgcc | gatcttt | | | 217 |
| <210> <211> <212> <213> | 19506 379 DNA Glycine max | x | | | | |
| tgtcaagccc | cccaacttgg | acaagtgttg | acttagtttt | ttctgcctag | gtgtctgggg | 60 |
| tggacatact | tttggctttg | caagtgagat | ctgatgagtc | atgtgagaaa | agccattgca | 120 |
| tttgaagact | ttccctttta | ctgaccgctg | gagagtgcct | taatgataga | catctcattt | 180 |
| tgtccattga | tgcaggcgtg | tgcatcacac | tcgcaatact | tttgcgtaca | tgtcactcgt | 240 |
| ggatgacgca | cacactggag | acgtgatgca | tgggtaagag | gggctgtggt | cgggtgcaga | 300 |
| aaattatggt | accactctat | cgccttacag | ctaccgaaga | gcatggctcc | tctttatatg | 360 |
| gagcatacgc | ttgatctgt | | | | | 379 |
| <210> <211> <212> <213> | 19507 396 DNA Glycine max | ς | | | | |
| <223> <400> | unsure at a 19507 | all n locati | ions | | | |
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| acatcctaat | agttaatggt | ttagtggttg | ctaaacaaca | aaagtttagc | aagaataacc | 120 |
| taaaggctat | gaaggtgtcc | ttggaacaaa | gcttgaagga | aactccaaaa | gtgggcgtga | 180 |
| aagccataaa | caagagaaaa | caaaaagagg | ttggccaagt | ccattgttga | tgaagacctt | 240 |
| ggtctgaaca | actttggcat | ttcaaggaaa | attanaaaga | tcatggttga | aaaacaaaag | 300 |

| aagaacaaaa | ccagaaaaca | atctcatagg | gatgaagaag | aaaaggaaat | ataggcaaag | 360 |
|----------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| aagcatagga | ccaaaagaag | gcatgagtct | attacg | | • | 396 |
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| <223> <400> | unsure at a | all n locat: | ions | | | |
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| tcctagtttg | gacctttccc | tgttgctctc | taaggaccct | ttgaaccttt | ttatcaagtt | 120 |
| caaaggtctt | caaaggctta | tcttcactag | caacagtctt | ggccagtttc | ttctgcctcc | 180 |
| tttgtaggat | ttcatcttga | tttacttaag | gaatagggtt | gttccttcta | gtgtcaactt | 240 |
| tcattgcctt | taggttaaaa | gcactgaagt | gttgctatct | ggcaatgaaa | gttcccttgc | 300 |
| tgttgatgga | atgcttggag | tcctcatcta | gtctttcaaa | ggcatggaat | acctagtgtt | 360 |
| cctagaggaa | gttgttgatg | agggtggcag | agtagatatc | ttccactcct | cccaaggcct | 420 |
| tcatg | | | | | | 425 |
| <210> <211> <212> <213> | 19509 382 DNA Glycine max | K | | | | |
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| tttatgactc | tcgtgctaat | tctatcacag | tgtaaatatt | cttatatcag | tttgtccaag | 120 |
| aacataatca | tgtgcttaat | atcattcatt | aacggtcacg | gatctatcgc | actattaacg | 180 |
| aactatggat | atcagaccta | ttgttaatta | cattttgacc | gtataaatat | cttactgatt | 240 |
| caagcagata | gacgcttaat | acgaagttta | attaactcta | atggctcctg | gaggacccga | 300 |
| atttgaacat | gaattacatt | gatcttgcac | catgttttaa | acacctctga | cacactatcc | 360 |
| agctattatg | catcatgctc | ag | | | | 382 |
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| agccttgatc | aaaggtgaac cacgtaaatc | tgtgtgttct | ttctcatctc | tctccccctt | 120 |
| tcaatttgct | gcaaaatctg tgtgtatggc | atttctgttc | tgttgcatct | actgttgttg | 180 |
| ttcttgattg | ttcttcatca cttccataac | aagaatgata | ttataacaca | ccaggtaaga | 240 |
| ggaaagaaga | agaaacaaaa taggattata | aaatagacgc | aaatgattat | aatgaaaaaa | 300 |
| aaaagtgttt | tacctatgat gaggatgaag | gcaacttaga | ttgagtttat | gaaaaaggag | 360 |
| aggatgtggt | ggtttccata tttacctagc | aaaaaaaaa | aaaaaaaag | aatagttcat | 420 |
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| cttcataatg | gggccatgtc tgagttgtac | accgtgcaga | tactgcaggg | ggtataatcc | 120 |
| tatgcccgtc | tacagtgggc ttattgataa | ttgagccgga | ctcttcctag | atatagccga | 180 |
| ccaactgtga | catatgatat a | | | | 201 |
| | | | | | |
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| <211> | 387 | | | | |
| <212> | DNA | | | | |
| <213> | Glycine max | | | | |
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| tctgatcata | tactacaacg tccctagctt | aatttgcaga | ataatgttac | atatatatgg | 120 |
| catcctcaaa | ccacatatgg catatactat | ttattttcat | cttttacatg | atgtatatct | 180 |
| ggaccgaaag | agacacgggt gacgggtcca | ccccatcatg | aaccatttat | tttttataat | 240 |

| gttgaaagaa | ttggcatata | tagtagatta | attctaattg | gtttgaccat | gatcaaggta | 300 |
|-------------------------|------------------------------------|-------------------|------------|------------|------------|-----|
| tctgaatttt | ttgtacagaa | tgaatatttt | agtactgtca | ttgcaagaga | ttattncgta | 360 |
| tgagaactat | tcataaacat | ccatcat | | | | 387 |
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| <400> | 19513 | all ii locac. | TORS | | | |
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| agttttaaat | gatctttgga | tagattcttg | aaataacttc | atagattttt | agaagcattt | 120 |
| gagagaataa | ataagttaca | taaaatttca | tgataagatt | ttttaagaga | cttagggtca | 180 |
| ttgaaattta | ccacatcttc | ctagtctgat | ttggatcctt | caaaagttgt | gtctgtcatc | 240 |
| aagcatatgt | tggcttcttc | atcagacgag | gtgtcgtcta | ggtcttcaca | tgtgctcata | 300 |
| agccccttat | tttctttggt | cttaaagaat | ttcttcttat | cttgactttt | ctcanaatct | 360 |
| ggacattcat | atttgaagtg | tctaggcttt | ttgcattcat | agcaaactat | gaagcttttg | 420 |
| tttttattcc | tttttt | | | | | 437 |
| <210> <211> <212> <213> | 19514 368 DNA Glycine max | к | | | | |
| <223> <400> | unsure at a | all n locati | ions | | | |
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| aagcatactg | tttatgtcaa | aaacttgaaa | agagacagta | gcaatagcag | aacaagatgc | 120 |
| ccaagagaag | tgtacagcct | cacgttccat | caagatgagg | tgggggatca | agtcctggga | 180 |
| gccatgtatg | tcgctgttgc | gccccaacac | aagtttgcct | ataacgtgga | gtggcagcag | 240 |
| caccatcccc | aagccacgat | aatcttccag | ctggtgaacc | tctatccaag | ctcttggtgt | 300 |
| tgaataaaga | gccgggctga | atggagaagc | ttcttcctat | tgctacgggc | attggtntct | 360 |
| gattatct | | | | | | 368 |

| <210 <211 <212 <213 | > > | 19515 424 DNA Glycine max | × | | | | |
|------------------------------|-------------|---|-------------------|------------|------------|--------------------|------|
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| atgc | tggtgt | aagaaccgtg | ttttcatttt | gtttcttatg | ttgngatcct | tgatcatcaa | 60 |
| tcat | ttcttt | ctggaatagc | tgatgttcgt | gatcggcaca | gggatatgcg | acttgatgtt | 120 |
| gata | acatgt | cttatgaggt | aaaatctcat | ttctacatct | ggatggaaat | attaatatat | 180 |
| ctat | acaacc | atagtgtcag | tttatccctt | atatttcaag | tcagtgcttc | gctttagaga | 240 |
| gttt | ctcatg | tgacattcgt | gactataaca | ggagttgttg | gctctggaag | agcgcattgg | 300 |
| aaat | gtgagt | actggattga | gtgaggaaac | tgtattgaaa | cacttgaaac | agagaaagca | 360 |
| ctcg | gctgan | aaagggcctc | agattgatgc | agaaccctgt | tgtgcttgtc | aggtaaacct | 420 |
| gact | | | | | | | 424 |
| <210 <211 <212 <213 | > > > | 19516 363 DNA Glycine max 19516 | × | | · | | |
| agct | tctatg | gaggctgaat | ctttgagctt | caatgaggtc | cttcaatgat | gattttcaac | - 60 |
| catg | gagatg | cagcggaaga | taaaggagaa | gaggtgagag | gaggcgtcat | ccactatgga | 120 |
| ataa | gccatg | gaaggagaag | cttcaccacc | aagagagtgc | cttggataag | aagcttaga g | 180 |
| agga | agcttc | aatggaggaa | aagaatgaga | aggagagaga | gggggaggga | ggcacgaaat | 240 |
| tgaa | ggagaa | aaagagagag | aagttgaact | ttgaagtgtg | tctcactagt | ttcacattcg | 300 |
| tcaa | aattat | gacaagtgtt | acacatgttt | caattatagc | ctaggtcatt | aactaaatg a | 360 |
| aag | | | | | | | 363 |
| <210 <211 <212 <213 | > > > | 19517 433 DNA Glycine max | K all n locat: | ions | | | |
| ~443 | _ | unpure at a | .טטטע זו בבג | T (1110) | | | |

| <400> | 19517 | | | | | |
|----------------------------------|------------------------------------|-------------------|------------|------------|------------|-----|
| tgagcatgtc | aacatgaagc | atatntctta | actgtcgtta | actgatttac | taaaaggact | 60 |
| acccaagatt | tgttggaaaa | ctcatcttct | ttgtgaagca | tgttataaag | ggaaacaaat | 120 |
| caaaactacc | tttaaatccg | gagatattgt | ttccactacc | agacctttgc | aattgttaca | 180 |
| tatggaccta | tttggaccta | caagaacttt | gagtctaaga | ggaaagaaat | atggctttgt | 240 |
| catagttgat | gactattcta | gatacatgtt | ggtatagaga | aaacggttat | aactgtctgt | 300 |
| aatttattaa | atctataagg | taattgatta | ttgtaacaaa | gttaccaatt | agattatcta | 360 |
| agtaatcaat | taaagtgttc | atccaatata | tggaaaacaa | ctcaagaaca | atgtaatcaa | 420 |
| ttatatgacc | tga | | | | | 433 |
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| <400> | 19518 | | | | | 0. |
| | | | | gggagagaaa | | 60 |
| gcagcacgaa | attgaaggaa | gaaaaaggga | gagaagttga | actttgagtt | gtgtctcaca | 120 |
| agactctcat | tcatcaaagt | tacaacaagt | gttacacatg | cttctattta | tagactaggt | 180 |
| agcttccttg | agaagcttgt | ttgagaaaac | ttccttgaga | agctagagct | tagctacaca | 240 |
| cacccctctc | ataactaagc | tcacctcctt | gagaagcttc | cttaagaaga | ttcctaaaga | 300 |
| tgtttgagct | tagctacaca | tacctctcta | atagctaagc | tcacctcctt | gagatgagaa | 360 |
| gctaaagctt | agctacacac | cccctataat | agctaagctc | a | | 401 |
| <210> <211> <212> <213> | 19519 432 DNA Glycine max | x all n locat: | ions | | | |
| <223> <400> | 19519 | art II TUCAC. | TOITO | | | |
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| catgagatgt | gaaattggaa | agatgctaat | ctacatagtt | catacttttt | tataaatgtt | 120 |

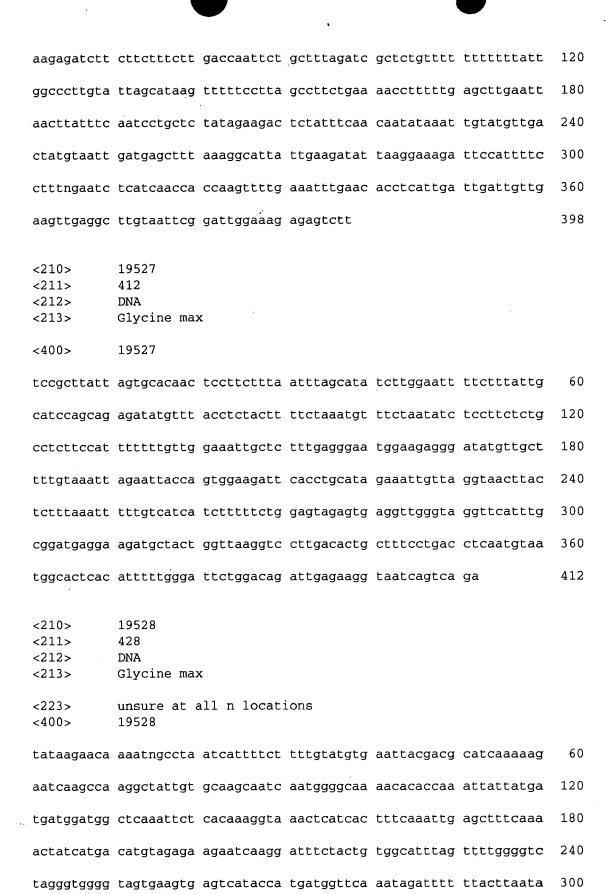
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| aggtatcaac | caagaaaaga | aaatacagcc | aaaaccaaag | cagagagaca | attagcaata | 240 |
|----------------------------------|------------------------------------|------------|------------|------------|------------|-----|
| aaactaacta | ataaaactaa | ggagtaagga | accaccaaaa | ttattcaggg | tctctgtcct | 300 |
| gtgttgtaca | tacattaaat | aaatggcaag | gttctaaaac | cactatgcat | aatctaacat | 360 |
| anaataaatt | atataaataa | atgcaggtac | cttaaatttg | atatggccaa | gccaagcttt | 420 |
| gaacataatc | at | | | | | 432 |
| <210> <211> <212> <213> | 19520 400 DNA Glycine max | ĸ | | | | |
| <400> | 19520 | | | | | |
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| ttatgatcgg | gtcaactaca | tcgtaactat | ctttgggaag | acccaaaaga | agccatcatc | 120 |
| tgagctaaac | atatggaaga | aaaggtcaat | attctttgat | cttcgatact | ggtccgatct | 180 |
| tgatgttaga | cattgtatag | acgtgatgca | tgtggagaaa | aatgtctgcg | atagtttaat | 240 |
| tggcactctt | cttaacatta | aaggctagac | aaaggatggt | ttgaagtgtc | gtcaatactt | 300 |
| ggttgagatg | ggtatacgag | agcagttgca | tccgatctca | caaggtccac | gaacgtatct | 360 |
| gcccccagca | tgttagataa | tgtcaacaaa | agagaagtga | | | 400 |
| <210> <211> <212> <213> | 19521 434 DNA Glycine max | ς. | | | | |
| | | | | | | |
| tgttgtcttc | gtttagtttc | ttggttgtag | atatcttttt | atacaatccc | accttgtgat | 60 |
| atccaccaga | attgttcgca | tatgaagcat | acatcaatgg | ggggtataat | tccaacctta | 120 |
| ttgaaaaatt | ttatttgcat | aatatcactt | gtgctcaatt | tcttgtgacc | aagtaacatt | 180 |
| gcacaatcca | tttcatccaa | catgtcatgg | ttgtggtcaa | agtcaacata | tgtaacatac | 240 |
| caatgatcgt | tactaaagtt | aacgtgaaca | tgaaacctag | cattgcaacc | accttgtatc | 300 |
| atttttcct | cacgcttcct | agttttcaat | gtcaaaccac | tatcatttca | atatccaaca | 360 |
| | | | | | | |

tgagagcaaa taaaagtttg ttgtaatgtt ttccctataa tgtttctcat aacaatactt 420

| ttacggacaa | agaa | | | | | 434 |
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| <210> <211> <212> <213> | 19522 384 DNA Glycine max | : | | | | |
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| aacatcgtgc | actggttaaa | ggaaaaaaat | tattcaagat | cacttggtag | cgtctttggc | 120 |
| attacgtggt | atatggtttc | tttgactact | ttataatgaa | attataacgc | tcattttgat | 180 |
| tctatttact | tgattaaggt | tttcttctaa | aataggcaca | gtctgtttgc | tattgattct | 240 |
| cttgtttgga | aaagggactc | ttgatttaat | cttgaaatca | taggaatttg | acttctctaa | 300 |
| tatgtgtaac | ttactttaag | ctcatggttg | cagaggaaga | acaatacaaa | tagatcagtg | 360 |
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| | ataatgattn t | tagatcaact | aatacgtttt | cataggaaaa | tttgaacacg | 60 |
| | cttaattcag (| | | | _ | 120 |
| | aagtttaatt a | | | | | 180 |
| agcttaatag | gttatatata t | tataaaaatg | tacatcatat | cgatggaaaa | ttcatattta | 240 |
| attttattct | gcttctgcct o | cagctccatc | tattactctt | gctcaacaag | aaaatattag | 300 |
| cagcttagat | attgagcatt t | cacaaagtca | tcttggacac | aagctttctg | gtcagaaata | 360 |
| tttactagta | ttggatgata t | atggggtga | tgatcgtgca | caatggatag | tgttgaaaga | 420 |
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|--------------|-------------------------------------|--|--------------|------------|------------|--------------------|-----|
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| rus e ser | <223> <400> | unsure at a | all n locat: | ions | | | |
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| | agatatctta | tgaagggggg | gttgaattaa | gatattccaa | actacttccc | caattaaaat | 120 |
| | ctatttcact | ttttattcga | gttataaatt | cccttaataa | tgaacttctt | aaatattgat | 180 |
| | tcaaataaaa | caatttgaat | atgaatataa | agcaataáta | aacaaaggag | attaagggaa | 240 |
| | gagaaaatgc | aaactcagat | ttatactggt | tcggccacac | ccttgtgcct | acgtccagtc | 300 |
| | cccaagcaac | ccgctngaga | gttccactat | cttgtaaatt | ccttttacaa | gttctaaaca | 360 |
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| | ctattttcag | attgggaatg | cctctaacag | cacctttgtc | aatgattttc | ttcatgcctc | 120 |
| | ttaagtgcag | atgtccaaat | ctttgatgcc | atatattgac | ttcatcttct | ttggagact a | 180 |
| | gacatgtgga | ggagtaactg | gtttcttgag | gtgtccatag | gtaacagttg | tcctttgatc | 240 |
| | tgctgccctt | cattaggact | tcactcttct | catttgtcac | caagcattct | gactttgtg a | 300 |
| | agtttacatt | gagtccttca | tcacacaact | gactgatgct | gatcaagttc | gcagtcagt c | 360 |
| | ccttcaccag | cagtactttg | ttcagactan | gaagtccttc | atggactagc | tntcccattc | 420 |
| | cagtgatc | | | | | | 428 |
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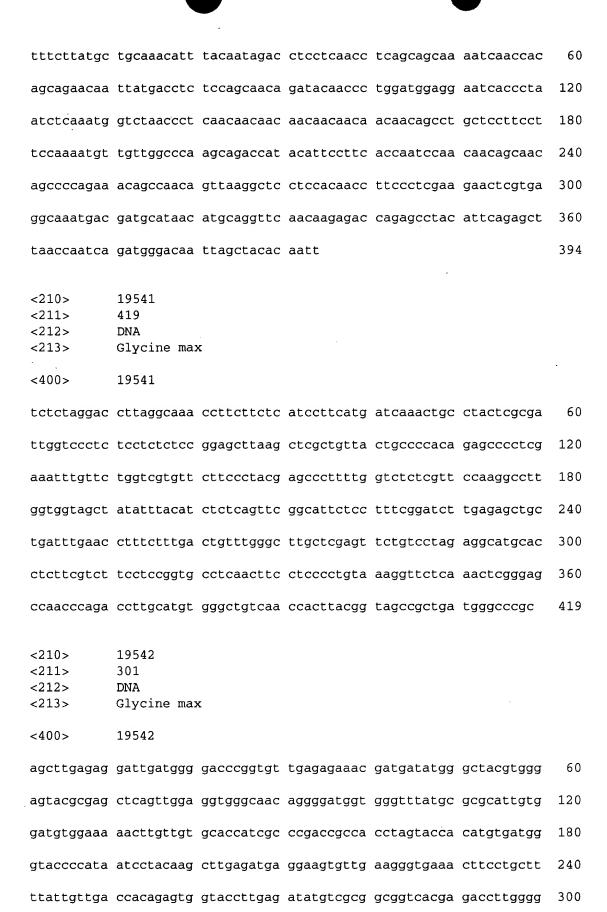
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|----------------------------------|------------------------------------|-----------|------------|------------|------------|-----|
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| caacaata | | | | | | 428 |
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| gtggatggcg | ccgcctctta cc | tcttctcc | tttgtctttc | gctgcatctc | catggtggaa | 120 |
| aatcatcatt | aaaggacctc at | tgaagctc | aaagatccag | cctccataga | agccccacaa | 180 |
| gcaagcttcc | atcaagtggt aa | tcagagca | caagagcttc | aattaggtgc | tccttaaacc | 240 |
| tccattaatt | tttttgcttt ac | cttctctt | ccattattgg | ttcttcattn | tttctccatg | 300 |
| tatctcctca | catgtcttgg tc | tanatgtt | gttaacatga | ttctttagag | tttcaaccaa | 360 |
| ttaaacttgt | tatagaagct ag | atttgant | ttctat | | | 396 |
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| ttntttttgc | tttaccttct ctt | tccattgt | tgtttcttca | tttttttctc | catgtatctc | 120 |
| ctcacatgtc | ttgtgctaaa tgt | ttttaac a | atgattcttt | agagtttcca | ccgattaaac | 180 |
| ttgctatata | agctagattt gat | tttctat q | ggttcaaatt | tcttgttttt | gttcttgaac | 240 |
| catgaattgt | gttgagttta ggt | tcctttg a | agttttgtct | tgttattttt | tgtggctgaa | 300 |
| acctaaacca | taaaattcat aca | aaaatat t | taaagtagaa | taaaagctca | taaatctaga | 360 |
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| acat | | | | | | 424 |

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| aatttgaa | aa gtcatgaccc ttcaaaatat aactgtgtaa tcgattacca gtgaagaatt | 120 |
| tcagaaaa | ag ctttttgaaa agacacattt cttgaaatca ttntgaaaag gcacgaaggg | 180 |
| cctatata | ta tgtgtgtctg acttcgaaaa gcaagagaga gattctaaga gaacttaatt | 240 |
| gtcaaatg | ct ctctcaacaa ctcttgggca aacacttgca aatctattga gaattcatct | 300 |
| aggaactt | ca aattgtatta tcatctctaa aag | 333 |
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| tgcttccaa | aa gtttcatggc cttgcaggtg aagacccgca caaacatctg aaagaattcc | 120 |
| atattgtct | cg ctccaccatg aaacccccgg atgtccagga ggatcacata tttttgaagg | 180 |
| cttttcctc | ca ttctttagag cgagtggcaa aggacttgct ttattacctt gctccacgat | 240 |
| ccatcacaa | ag ctgggatgac ctcaaaagag tattcttaga aaaaaaattt cctgcttcca | 300 |
| ggaccacga | ac catcagaaag gatatttcag gcattagaaa actcagtgga gagaacttat | 360 |
| atgaatact | g ggagagattt aagaagctat atgccagttg cccgcaccac cagatttctg | 420 |
| agcagcttc | et tetecaata | 439 |
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| <400> | 19533 | |
| cagtttttg | g tagacctaca ctagcggtga tgcactacga ggttgatcct aatgatcctg | 60 |

| atccactcaa | ggatttatta | cggctacgtg | atcaactttt | gagcaagctg | aaaagtaatt | 120 |
|----------------------------------|------------------------------------|------------|------------|------------|------------|-----|
| · | | | | gaaaataaga | | 180 |
| | | • | | | | 240 |
| ttaacgetgg | ayatatyatt | ctagttaagt | tacaycctca | catgaaacaa | ccaycyyccc | |
| tgaggaagca | tcagaagcta | tgcatgcgct | attttggtgc | gtttatagtg | attgaaaaaa | 300 |
| ttggtacgat | tgcatataaa | gaacaactgc | ctgagtc | | | 337 |
| <210> <211> <212> <213> | 19534 441 DNA Glycine max | κ | | | | |
| <400> | 19534 | | | | | |
| cttgcgattt | atccagctgg | aatccataaa | tgatgaggtg | atattgaaag | cttgctgttt | 60 |
| atgcaaaaaa | atggaatcca | gaaatgaagt | gacattggaa | gctactctca | atattgcaca | 120 |
| atgtttcata | gcctctcttt | gtcacaatag | ttcggcgatg | tgaatcaatc | atatcaaact | 180 |
| ataagatcga | catatcctga | aaactagccg | aaccatagcc | tgcatttttg | aattcgaatc | 240 |
| tcaaatttgg | aaatgggaaa | gggtggaagg | aagaagcaac | agataccttg | cacgctcttg | 300 |
| ctgatggaag | attgaggaag | ctaacacgtg | ttaggaagaa | ataaaaaatt | ggaaaaaatg | 360 |
| gatagcacta | acacgggaaa | tcggttgttg | tttattacta | acatacagac | tagtactatg | 420 |
| atatcataat | agtctttagt | С | | | | 441 |
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| agctttttgg | agtagaaaca | tgggaccaac | tcattttatt | tcacaaagga | agtcgtatct | 60 |
| agtcaaggtc | tgagagacca | tacaagtttc | ctaacgattt | ctaattatgt | gggccattaa | 120 |
| gactatcata | tgctgacaat | agccgagaag | cccatgaatc | tcttcggggg | cggagtaggt | 180 |
| gtctgccatc | gccttggcct | tggctaacaa | tcggggaagt | tcttgactcc | cgttcaaggt | 240 |
| aagagcatac | cgatccatcc | acatggatgc | ctcttggtgt | aaagagtcga | tcaccctttc | 300 |
| tctagcctct | ttttgcgcgt | atacttgggc | atattcgtcc | gcaatcctat | gctcctgggc | 360 |

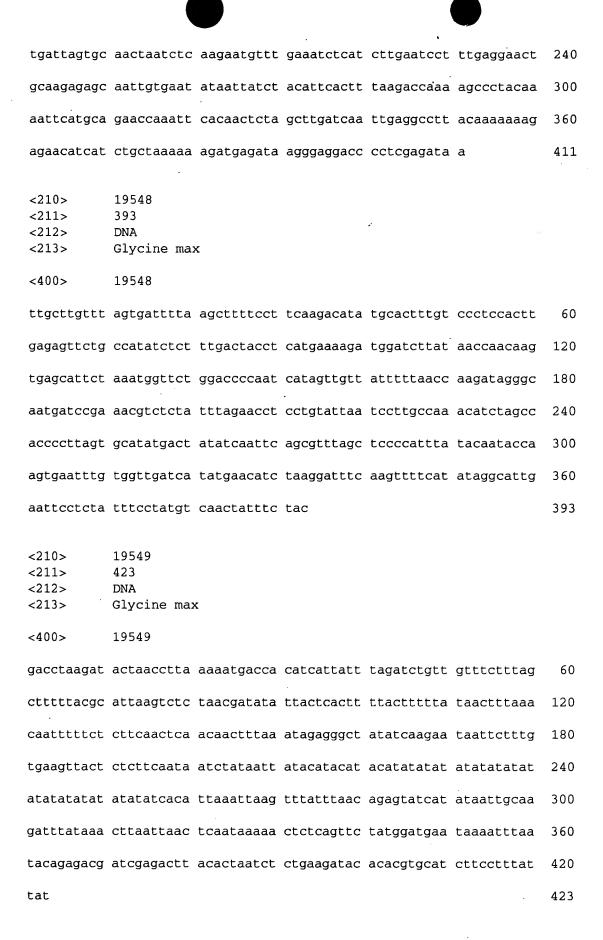
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|-------------------------|---|-------|
| <210> <211> <212> <213> | 19536 441 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19536 | |
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| gatcgtacag | g cgggtgaacc ataagcggaa gtttcttttg gtgaggtagc catggaaaa | g 120 |
| catagcgttt | ggaatgattt cgtatatctc agaaggctat tgggaaatgc tggttaaaa | c 180 |
| acgaatgcca | a agcagatata aattttaatg aagaatgtat atgggcgtgt gacgcaacg | g 240 |
| tcgaatttgc | c tttgcggtga acgtgctatt aatgttaagt gattcgtttg ggcacgttc | a 300 |
| gattgcagta | a gctgctataa tttctctagc agacaaatgc ccatcttgcc cctcagttt | 360 |
| tcaaactgat | tagcatccaa agcctttgtg aaaatatctg ctattngctg ctcagtgtc | a 420 |
| acatgctcta | gtgtgatcac t | 441 |
| <210><211><212><213> | 19537 401 DNA Glycine max | |
| <400> | 19537 | |
| agcttctaag | aaagcttctc aaggaagcta cctagtctat aaatagaagc atgtgtaaca | 60 |
| cttgttgtaa | ctttgatgaa taagagtett gtgagacata etteaaagtt eeacttetet | 120 |
| ccctctttta | ttccttcaat ttcatgctcc ccctctctct ttctctccct ctttcttttc | 180 |
| ctccattgaa | acatecttee aagettetta tecaaggete atettggtgg tgaageteet | 240 |
| tcttccatgg | cttattccct agtggatggc gcctcctctc acctcttctc ctttgtcttc | 300 |
| cgctgcatct | ccatggtgga aaatcaccat taaaggacct cattgaagct catagatcca | 360 |
| gcctccatag | aagccccaca agccagcttc catcaagtgg g | 401 |
| <210> <211> <212> | 19538 436 DNA | |

| <213> | Glycine ma | x | | | | |
|----------------|--------------|--------------|------------|------------|------------|-----|
| <400> | 19538 | | | | | |
| gcttgtcatg | atccgtctct | ttggtgcaaa | attgatatgt | ccttacttaa | cagcttataa | 60 |
| ttttttaaca | taccaaataa | tcagccatga | gcttacaagc | gtacaagtgg | aaaaattact | 120 |
| caactcttaa | agtatgttct | acgtctgagt | aatggaaata | catattgctt | agtatttaac | 180 |
| tacaatgttt | acttgactga | tgagtagttc | atcatagata | ttgaaaggta | acattgtctc | 240 |
| ttattcttaa | ttacccctta | atttgtacat | gcattattaa | ataacctttt | aaaacaaaaa | 300 |
| tacttcatca | atattagtgc | tcaagtctaa | attaaatgcc | acgtataata | tttatataaa | 360 |
| agttgttgtc | atatgggatt | gataagcgtg | cgtgtgcctc | gttatgttag | gaccccaaat | 420 |
| ttgaaacaac | tagttc | | | | | 436 |
| | | | | | | |
| <210> <211> | 19539 · 496 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | κ . | | | | |
| <223> | unsure at a | all n locati | ions | | | |
| <400> | 19539 | | | | | |
| gatgatcgnc | natgaaccct | ttggatgaac | gcgatactat | agaacactca | agtgggaccg | 60 |
| tggtcccaga | ctaataatca | gaccgacgat | actattggga | ccgtggtccc | agactaataa | 120 |
| tcagaccgac | gatacgagtg | ggaccgtggt | cccagtctga | ttatcagacc | gacgatacaa | 180 |
| gtggaacagt | gggcccagag | agaatattca | ggccagttat | gctttctggc | ctgtaacaaa | 240 |
| ggacattaag | taaagacaga | taaacgtaga | ctaaaacgtg | gtcgcatcag | ggtgctggct | 300 |
| tttcaagttc | cttaagaatg | gcctcaattt | tctctataca | ctcagttgga | acacgagacc | 360 |
| tgtccaggtt | aagcaccatt | ttatcgccct | tatacaatac | tgtcgctcca | ggagcaaact | 420 |
| gatgtcgaga | gcttaaacta | gttcttgatg | cagatgacgt | tttaagcaca | gaagttaaaa | 480 |
| gagtgataac | ttcttn | | | | | 496 |
| 0.4.6 | | | | | | |
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| <211><212> | DNA | | | | | |
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| <400> | 19540 | | | | | |



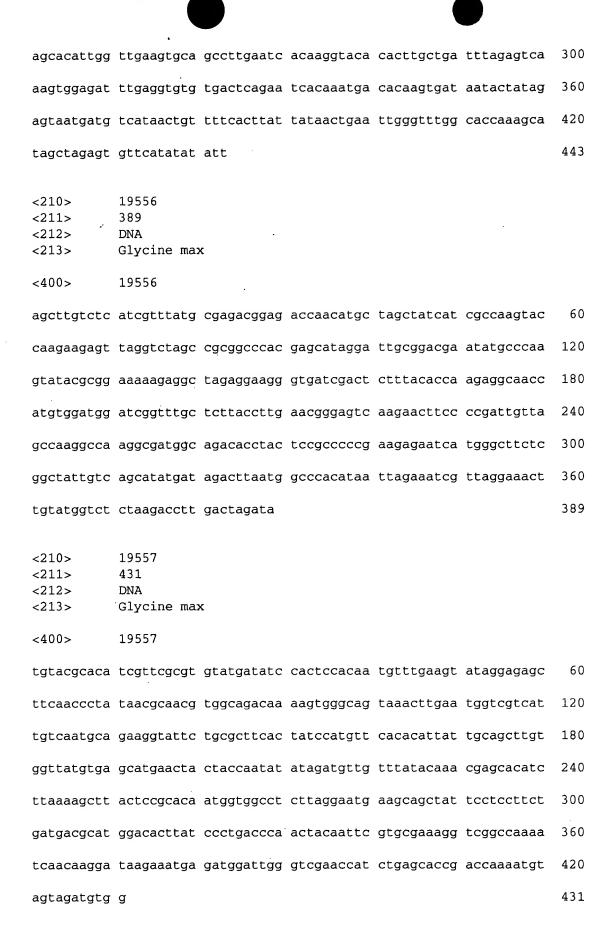
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|----------------------------------|------------------------------------|-------------|------------|------------|------------|-----|
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| <223> <400> | unsure at a | all n locat | ions | | | e. |
| acaaactcaa | gcttaggaac | ccaagctctt | agcttcaatg | caaggaaaca | tatttatgcc | 60 |
| taataaccta | actnttggtt | gtggaagtag | gaaggcatga | aaattatgac | ttgcttgtga | 120 |
| gagtttntac | tcgaatttgg | gctgccccat | gaggggtaat | ttgcacctaa | gtagtgtgga | 180 |
| aaatacttta | caatggtatg | taaatatgtg | tgtaaatata | cgnggcatgg | aaaacacctc | 240 |
| tcaatggtgt | gtatatatgt | gaacatatgg | catgaaattc | cttgcaaagc | gtgaatgagt | 300 |
| atcttcctaa | atgaatagag | gtcgcttcct | aaatgaatgt | atgatggcat | ggaattccct | 360 |
| ttttacatgc | aagtatgtgc | atgacgtaat | tagctttcca | atatgcatat | aaataaatgt | 420 |
| gagtgaaaca | atgaaagt | | | | | 438 |
| <210> <211> <212> <213> | 19544 388 DNA Glycine max | · · | · | | | |
| <400> | 19544 | | | | | |
| ttgcttgtga | aataaaagtg | tcaatatgtg | tagtgtatac | actggggcgt | cgaaaattta | 60 |
| aagaaaagaa | tċaacaagat | tgaaaggcta | atatatcctc | tataacaaaa | tcacaaccac | 120 |
| acaatattta | tgctccttat | aaagaatcct | aacgcctaag | gtacacactc | aacacaagaa | 180 |
| cacatcaatt | ttacaacaaa | ttcgcatcga | aacaccaatt | ggtccatcaa | acacactaaa | 240 |
| tccgtgatta | aaacaaaaca | acacatagtt | gaacttcata | aaacattcca | aaataaccca | 300 |
| taaattgatc | ctcgatgtag | tcgctcaagc | gttattcgct | agcaatgaca | ttactggtgt | 360 |
| tctctaaagc | tcctcttccg | attgctct | | | | 388 |
| <210> <211> <212> | 19545 396 DNA | | | | | |

| | | | | | • | |
|-------------------------|------------------------------------|------------|------------|------------|------------|-----|
| <213> | Glycine max | x | | • | | |
| <400> | 19545 | | | | | |
| tcaaggttat | gacttcatgt | tgctcatcct | atctctaata | tacacaccac | aaattctcct | 60 |
| catgttagcc | tttgcctttg | agttccaccc | atattttagt | gcaaaccaag | aacctgagaa | 120 |
| gattatactc | attctttacg | gctgggttag | gttgagtgga | ttggaagcaa | gaaaaaaata | 180 |
| ctccctccgt | tactatttac | aaaaggttgt | agtcgaattt | aagctaatta | agtattaaca | 240 |
| agtactaata | tcaattaata | agacattaag | aagtaacagt | aacaactata | aaaaatagta | 300 |
| tcaattatta | atcatcaatc | atgcgtcatc | aatatatata | ttatcttatg | tataaattat | 360 |
| aaaataagat | cttatcacat | aacaatgact | atgttc | | | 396 |
| <210> <211> <212> <213> | 19546 395 DNA Glycine max | ĸ | · | | | |
| <400> | 19546 | | | | | |
| ttgcttttca | ctttataaag | ggagagttta | gtatgaattt | tgatgatacc | atctgatgta | 60 |
| attctacccc | ccaagggtat | tggataaaaa | acttcaagaa | gattgagcca | gagatgtaag | 120 |
| agaaggtcct | aggattctca | tgagccttat | ggtagatttc | ggacccatgg | gctaagtatg | 180 |
| agctcactta | tctttgtaca | tattagatta | aggtttcatt | atttttgtgc | cttgtattta | 240 |
| gagctccata | atgtagatag | ggtaccctag | agatatagga | attttcaacc | cttgtatttt | 300 |
| aaggcaccta | gactagtctt | tgtattatgg | gtagttttgt | aatttcactt | gcattaagtg | 360 |
| aatatttgat | gtgtgtgttg | ggaaataaat | ttaat | | | 395 |
| <210> <211> <212> <213> | 19547 411 DNA Glycine max | ς. | | | | |
| <400> | 19547 | | | | | • |
| tgcataggat | aaattggagt | ggaagccaag | actgagtttt | ctgagacata | aacaacatgc | 60 |
| cctatttaac | aacttacctt | ttcaagaggc | caatctagat | tgcaccttct | ccacaacatc | 120 |
| atcaaagtca | aatttagtca | ctcagtcagt | tagaggtttg | aaacccaagt | acttcccaag | 180 |



| <210> <211> <212> <213> | 19550 51 DNA Glycine max | |
|-------------------------------------|---|-----|
| <400> | 19550 | |
| agcttgttga | tgtggaacta ttgcttgcat gtgggtgtgt ccgtcgcttg a | 51 |
| <210> <211> <212> <213> | 19551 188 DNA Glycine max | |
| <400> | 19551 | |
| tgtcacgcaa | gctttcatca ccaactagcc ttattgattt taactgcaca gcgactttaa | 60 |
| gggcccaatg | agtcctcgat gcttatatca cgaacttatg tacgctcaac cttggctgac | 120 |
| gatatcaatg | gtgatcggaa ccataagtag atatgaaata ctttcaacct tggccgagag | 180 |
| aggttact | · | 188 |
| <210> <211> <212> <213> <223> <400> | 19552 377 DNA Glycine max unsure at all n locations 19552 | |
| agctttcttg | gaaaatgtta ttgttttgaa ggcaatgctt agaggctttg agatggcctc | 60 |
| tggtctgaaa | atcaactatg ctaagagtca attcggaatt tttggagatt atgttaactg 1 | L20 |
| gtctcaagaa | gctgctcact ttctgaactg tagacagatg gagattccct tccactactt 1 | L80 |
| gggcatcccc | atttgggtca gatcctcaaa tcaggtggta tgggagcctt tgatcagcan 2 | 240 |
| atttgaagct | aaactcacta natggaacca gaaaagctta tctatggctg gcagggttaa | 300 |
| tctgataaat | tctattttga acgctntacc aatctatcta ttatccttct ttaagttacc 3 | 860 |
| ccaaagaata | gctgata | 377 |
| <210><211><212><213> | 19553 420 DNA Glycine may | |

| <223> <400> | unsure at 19553 | all n locat | ions | | | |
|-------------------------|------------------------------------|---------------------------------------|------------|------------|------------|-----|
| teenettett | cccacgttct | tgcccttgaa | acctcacctt | ggtctcacta | ctcttcacat | 60 |
| cacgatgttc | ggtgtccttg | aagaggtgaa | cctcatctcc | tctatcttct | tctcttcaaa | 120 |
| caatcacgcc | aaataaaacc | cctttgacaa | cgagtgctcg | tcgcgaagat | aaaaaaccct | 180 |
| taccgaaccg | gatgagtaaa | aataagacgc | acaaaggaaa | atgtatccta | tgcaataata | 240 |
| acggtggacc | taagttcgaa | tatcaaactt | agaagactag | ttgtgttccc | aaataatcaa | 300 |
| tttagcaaca | attacgcaaa | ttgagtttta | tcaaattaag | aaagattgtt | tttgatcaat | 360 |
| ttactcttta | acgaagagaa | gaattaacac | acaagaactg | tgtgagaaac | atttataatg | 420 |
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| agcttgatat | gaggaagtgt | tgaagggtga | aacttcctgc | ttttattgtt | gaccacagag | 60 |
| tggtacctgg | agatatgtcg | cgggggtcaa | gagaccttgg | ggacgtcaag | tggggttcta | 120 |
| ttgcccaaaa | ccaagcttga | ccaatcccga | cccaacccgg | gcatagtcga | tcagtgagaa | 180 |
| cctatgatgt | acctaaacag | gcgagctcct | ggcagtcaac | agataaaagg | agcaaagacc | 240 |
| acaaagcaag | gaggcttgtg | gtggctggcc | agttgtgaaa | cttgattgat | atgtgagata | 300 |
| tggtctctgg | taatcgaata | ccaagggtgg | gtaatcgatt | acaaggctta | aaaatg | 356 |
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| <400> | 19555 | | | | | |
| tgcgtcacaa | ttcattgtga | cagtcaaagt | gccattcact | tagcaaatca | ccaaatgtac | 60 |
| catgagagga | caaagctcat | agatgtgaaa | ctacacttca | tcagatatgt | gattgaatct | 120 |
| gagaaggtga | aggtagagaa | ggtttcaaca | taagacaacc | cgactaatat | gttcacaaag | 180 |
| atcctctcta | gtgtcaagtt | caagcactgc | ttggacttga | taaattgtga | agatgcctaa | 240 |



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|----------------------------------|--|-----|
| <223> <400> | unsure at all n locations 19558 | |
| agcttgtgct | ccaacactgc ataggaagtg atgatttcgt tgcttgcagt gtgaaccaac | 60 |
| accttgcgct | ctgagccagg gttctccacc agcctcacca caccgttctt gaaaacccaa | 120 |
| accccagaca | a tattettage ttagaaacaa gaactagagg cacaacaaat tgaagaataa | 180 |
| gaagaacaaa | a ttaataccaa gtgttttgag tgtttacttt tgagtgcttc tgagttagtg | 240 |
| tgcaatgcaa | a gatggggagg ggagtatggg gtattgatgt atatatagtg gtggtgaatt | 300 |
| gaagtcatgg | g gtaagatggg ttttaattnt tttttttata atgtagaaat aataaaaact | 360 |
| acatggtggt | gataaggaca gtatattgag atg | 393 |
| <210> <211> <212> <213> | 19559 460 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19559 | |
| cgcacactat | agacactcaa gcttgttgaa gggtaccttn tggcaaatcc tgattactga | 60 |
| atgtctttct | tgttattaat taagagettg getatggtge aaagtaataa getageettg | 120 |
| atggatctgc | c aagctcagca ccaaggcttt cttcttctat gatatctttt atttattttt | 180 |
| gttctttatt | aaaacatggg caattaatca tagattaata atttattagt tttttttta | 240 |
| actttcttca | a gcttttgact aatacatatg tcctttgcat tggctgcaaa ttgatgtgtc | 300 |
| gccgcttgca | a atttggttaa aaaataacct atccttagaa cagactaaga gacacagaga | 360 |
| tgttagggat | atttttattc aagttatgtg ctagggttta gagagaaaat attanaaaat | 420 |
| aaatgtccgg | g atagataatc cactgtcata tgggactgat | 460 |
| <210> <211> <212> <213> | 19560 393 DNA Glycine max | |

| <400> | 19560 | | | | | |
|-------------------------------|------------------------------------|-------------------|------------|------------|-------------|-----|
| agcttgtatg | gttaaagtct | cacgattgtc | acgtgctcat | gcaacaattg | ttagtcgtgg | 60 |
| ctatacgaga | catcttgcca | aacaaagtaa | ggttagcgat | aactcgcctg | tgctttttt | 120 |
| ttccatgcta | tatgtagcaa | agtcattgat | cctgtcaagt | ttgatgagtt | ggaaaatgag | 180 |
| gccgcaatta | tactgtgcca | gttggagatg | tattttcccc | ctgctttctt | tgacatcatg | 240 |
| attcacttga | ttgtgcatca | ggtcagagaa | atcaaatgtt | gtggtcctgt | ttatctacag | 300 |
| tggatgtacc | cgattgagtg | atacatgaag | atcttaaaag | ggtatacaaa | gaatctatat | 360 |
| cgtccagaag | catctattgg | tgagaggtac | att | | | 393 |
| <210> <211> <212> <213> | 19561 441 DNA Glycine max | ĸ | <i>;</i> · | | Aly. | |
| <223> <400> | unsure at a 19561 | all n locati | ions | | | |
| ntaagtntga | gggagtttga | taatggttgt | gcattcctat | atttgcaatc | tatttcacaa | 60 |
| caattatctc | acttttcctt | gaatcgtagc | taagtttgcc | cttgtttaag | ttagggaata | 120 |
| tatataatta | gttagatatt | ttcatacagt | taaaatttag | aaaacttatt | agcttttaca | 180 |
| tgtttttaca | gtaattcagt | cattttagtg | cacctggaag | gaaattgagg | gtttggaagt | 240 |
| gaaaattgat | cactcaatga | gtttgccaag | tagcttaact | aggaagccat | attagaa gaa | 300 |
| gacacgtggt | agctggtggc | taagcaagaa | gtctatctct | cttagcagat | ttctcttgag | 360 |
| gaagccatgt | caacagcatc | aaggctcgtt | gagtgatgca | gcctctttgg | aagaagaaaa | 420 |
| aaagtgatcc | catganaaat | t | | | | 441 |
| <210> <211> <212> <213> <223> | | c all n locati | .ons | | | |
| <400> | 19562 | | · | | | |
| | atccgatcat | | | | | 60 |
| caatacggat | atggctcccg | atcggaacca | gcttcagagt | atgactaagc | gagagcatga | 120 |

| gťccattaag | gaatatgeee | : aaagatggag | agatctcgca | gcccaagtcg | tacttcccat | 180 |
|-------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| gaaggaaagg | gagatgatca | caattatggt | agatacgtta | cccacgttct | actatgaaaa | 240 |
| gctgataggc | : tacatgccag | ctaactttgc | ggatctcgtc | ttcgccggag | aaaggattga | 300 |
| atccagacta | cgaaaaggca | agttcgaata | tgcttccaat | gtggccccca | acaacaatag | 360 |
| aagagccnca | gtaatgggcg | cgaggaaaaa | gga | | | 393 |
| <210> <211> <212> <213> <400> | 19563 440 DNA Glycine ma | x | | | | |
| ttccctttgg | catcatcaaa | acattcagct | tgatcctttg | tctacataga | tgactctcaa | 60 |
| aaagcacttt | ctaaaagata | agatcgaatc | aaaagtcact | aataagaaag | aacaagaaat | 120 |
| ggtatttata | attttaacaa | acaaaactga | tttcatcaat | tatcaaatgt | ggtaattgat | 180 |
| taatttgcta | aatttcctct | ttgttccgca | tttccaaaaa | catggtaatc | aattacaaat | 240 |
| tgtggtactt | gattatctcg | tttcacaaag | agcttctcaa | gcttccatgg | tttcgaaata | 300 |
| atcaattatt | ttgacacaaa | gagctattaa | agtttccaga | tgtgatggaa | gcttgcttgt | 360 |
| ggagcttcta | tggaggctgg | ttctttgagc | ttcaatgagg | tcctttaatg | gtggttttcc | 420 |
| accatggaga | tgcagcggaa | | | | | 440 |
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| | catatacaat | | | | | 120 |
| | tgttacttac | | | | | 180 |
| | tatggggtac | | | | | 240 |
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| tgttgtcatt | aaatcttaca | tgaattgtct | cttccatagt | caaggtttnt | gatttgtaca | 60 |
| ctctatatgc | cttggatgat | tcaaagtatc | caatattcca | aaatcacatt | ttgagtcaaa | 120 |
| ctttccaagg | ttatccttgg | ı tgctgaagat | gaaacactgg | catccaaatg | ggtgacagtc | 180 |
| ataaatgttg | ggcttatgtc | cctcccacaa | tacataggaa | gtctttttta | agattgacct | 240 |
| tatataaatt | atgttctgta | aataataagc | agtgattgct | gcttctgccc | ataagtgttt | 300 |
| cagagttaag | tagtcgttaa | gcattgttct | tgccatttcc | tgaagagatc | catttttcct | 360 |
| ctcaacaact | ccattctagt | gggatgttct | tggagtacac | aaattattat | aataccattc | 420 |
| tctt | | | | | | 424 |
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| agcttgtatg | ctaactggat | gcattggtta | acttggtaac | ccaactggcc | ttgaaccaaa | 60 |
| aatctgtacc | tgttgcaagg | gtctgtggtt | tgtgctcctc | tgctgaccac | catacagacc | 120 |
| tttgcccttc | catgcaacaa | cttggagcaa | ttgagcagcc | cgaagcttat | gctgctaata | 180 |
| tttacaatag | acctcctcaa | cctcagcagc | aaaatcaacc | acagcagaaa | aattatgacc | 240 |
| tct | | | | | | 243 |
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| tgagatgagg | aagtgtagaa | gggtgatact | tcttgctttt | attcgttgac | cacagagtgg | 60 |
| | | gggtcaggag | | | | 120 |
| | | | | | | |

| cccaaaacca | agcttgacca | atcccgaccc | aacccgggca | tagtcagcca | gtgagaacct | 180 |
|-------------------------------------|---|-------------|------------|------------|------------|-----|
| gtgatgtacc | taaacaggcg | aggtcctgac | agtcaacaga | taaaaggaac | aaagaccaca | 240 |
| aagcaaggag | gcttgtgtgg | tggctggcca | actgtggact | ttgattgata | tatgggatat | 300 |
| ggcctctggt | aatcgattac | caagggtggg | taatcgatta | caaggcttaa | aaatgaagac | 360 |
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| tga | | | | | | 423 |
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| catgcatgtt | ttcttactgc | cattattctt | tgcagccagt | tgaactttgg | agtggtaagc | 120 |
| agctatttag | cattatactg | cgcccacatg | ctaatatgag | agtctatgtg | aatcttactg | 180 |
| ttaaggagag | aaactacact | gaagacaaga | aaataaaaga | caagaaaata | gaatggaaaa | 240 |
| cattgtgccc | aaatgatggg | nttgtttatt | ttcgtaatag | cgagttgatc | tctggacaag | 300 |
| ttggaaaggt | tactttaggt | tagttactct | aaatttacct | tgtttatttc | atctcatatt | 360 |
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| catgggggtg | agaagaggtg | gaaggcgacg | ccgatgtcaa | cacgatcgat | ggaggaactc | 120 |
| ggggagttga | agaaagagga | ggcggcatct | caggagcact | catggttgtc | acattggcac | 180 |
| taaacaaggg | tgtagtgaca | gtggacggag | gaaccacgac | atgattagcc | gaaacgggca | 240 |

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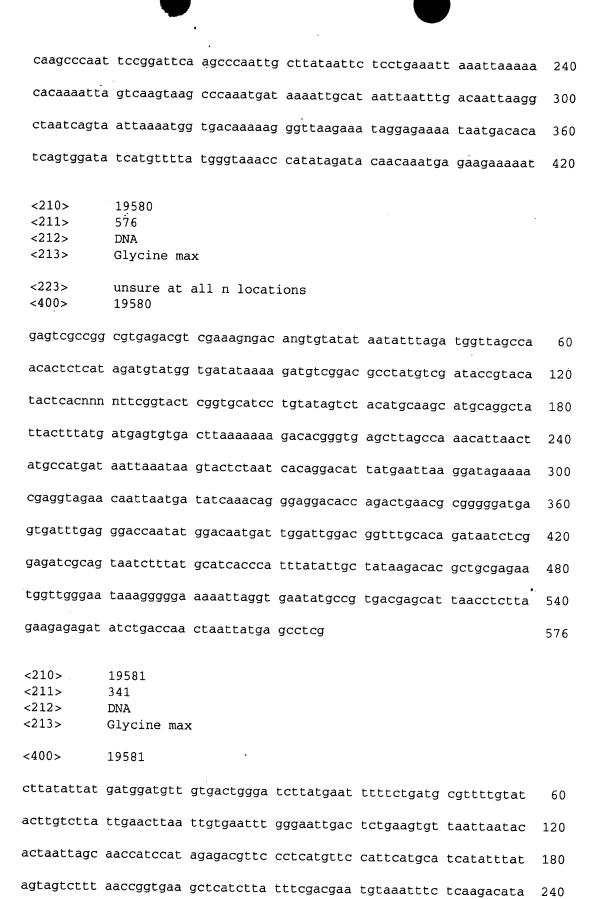
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| cctcatcacg | | | | | | 430 |
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| agtctacgga | ttgatagaat | atgttagatt | catatttact | ctctgacatg | aaccatgctg | 120 |
| aaagtattga | ttcgtgatct | atagcatcgc | ctattattat | attgatgtta | aatgaagtca | 180 |
| cgtgaaccgt | acttattcct | ttttatgata | gatcctgtga | agctattaga | ttgatccttc | 240 |
| gctcctatga | cctataccat | acgttataca | atgtcgagtg | acacatatag | ggatgttaca | 300 |
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| tcatttgct | ccaaagtttc a | tggccttgc | aggtgaagac | ccgcacaaac | atttgaaaga | 120 |
| atttcacati | gtctgctcca c | catgaaacc | cccagatgtc | caagaggatc | acatatttct | 180 |
| gaaggcttt | cctcattcat t | agagggagt | ggcaaaggac | tggctgtatt | accttgctcc | 240 |
| aaggtccato | acgagctggg a | tgaccttaa | gagagtattc | ttagaaaana | gtttccctgc | 300 |
| ttccaggaco | acagccatca g | gaaggatat | ctcacgtatt | agacaactca | gtggagagag | 360 |
| cctgtatgag | tactgcgaga g | attta | | | | 386 |
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| | atactcctta to | | | | | 120 |
| | ttataccagg to | | | | | 180 |
| | agttcgagaa ct | | | | | 240 |
| | tctcgaagag aa | | | | | 300 |
| | gaagagaact ta | | | | | 360 |
| gatgactgac | atttaggaac cg | gtaggggat | ttatactcat | aa | | 402 |
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| gatgcaacat | ctgtggtggg gc | ccatgagt | tatgtaagtg | catatcccaa | cacgatgcat | 120 |
| ccaaagaatt | caactacatg gc | taatccat a | atcatcgagg | gttccatcaa | ggaggacctc | 180 |

tgcgatacaa tcagggagaa acttttctta aggccaaggt tggagatccc atcctgtgaa 240

| tacattcatt | aaagattgac | • | | | | 260 |
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| cttttgacaa | taccaataac | tatggctaac | actaactatt | tacttttaaa | caaaacaagc | 120 |
| catatatcct | ggtcaagaaa | acatatacta | ggagtaattt | tctttataaa | tgattatgaa | 180 |
| aatgaattat | ccgcatctcc | atgttcccct | cctatccatc | aaaaggaata | agacaaggaa | 240 |
| ttttgttggt | caaacaaata | tcaaaataac | aatgtcaaaa | caacaattat | gtacaatgat | 300 |
| ttataattta | gtccccacac | catctataga | aaaaattaaa | caccttggac | ctagtaatca | 360 |
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| atgttgcaat | gagagcaaaa (| cggtttatcc | ttgcgagtag | gttttccctt | ggcattggat | 120 |
| tgggttacaa | cacaagccat t | ttcagcagga | ggagcataat | tgggttgaac | accaacaagc | 180 |
| ctatgagttt | catcttgaag a | aaataatgag | aaggtatcat | caatattcgg | agtcggtttc | 240 |
| atcaacaata | tttgacctca a | agcatgggca | aaactctctt | tgacccccat | cagaaatgac | 300 |
| atgacaaatt | ccnctttgat o | ggaagcaaga | agtggagcaa | ccctgccaca | attacaacta | 360 |
| tgattggcct | tgagttcacc o | cagcttagcc | ca | | | 392 |
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| <223> <400> | unsure at all n locations 19577 | |
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| cagagatato | c ttggcttgag ttactctcac ttggaaggct tcaatttaac ttcataatg | c 120 |
| tccttgttgg | g ccttttaaaa ctctaaagaa tatcgttgaa gtggatccca ctcaagcga | c 180 |
| aaattttttg | g aggaagcaaa caçaaatgat agttgtcatg tcatgggagc tagcaaggt | c 240 |
| attctgaggg | g aactttctga tgaacattct ggacttaaat cagaatggag attctaatg | t 300 |
| tagcttatga | a aggattcata ctgaagtaga teettttgat gaagtteaae agatgtata | t 360 |
| ctttctgaag | g tageteaget teateaacet taacacacet teataacaca nagtattet | a 420 |
| a | | 421 |
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| attgatctca | tagaacaact tgagaagttt gggaaagagt tttctcaaga tttgattcta | a 120 |
| caatcacttt | ctaattcatt ttcacaattt agtgtgaatt tcaacatgaa taagatgagt | 180 |
| tgtgacttgc | atgagatget aaatttgeta attgattatg agaateaaat tgettetgag | 240 |
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| aaaaaggaag | catcat | 316 |
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| gattggatga | aataaaatct agataagata agataagatt ggatgaaata aaatctagat | 120 |
| aaaataaaat | atggataaga taagatttga taaaagaaaa ttgtttgctc tcttcaagtc | 180 |



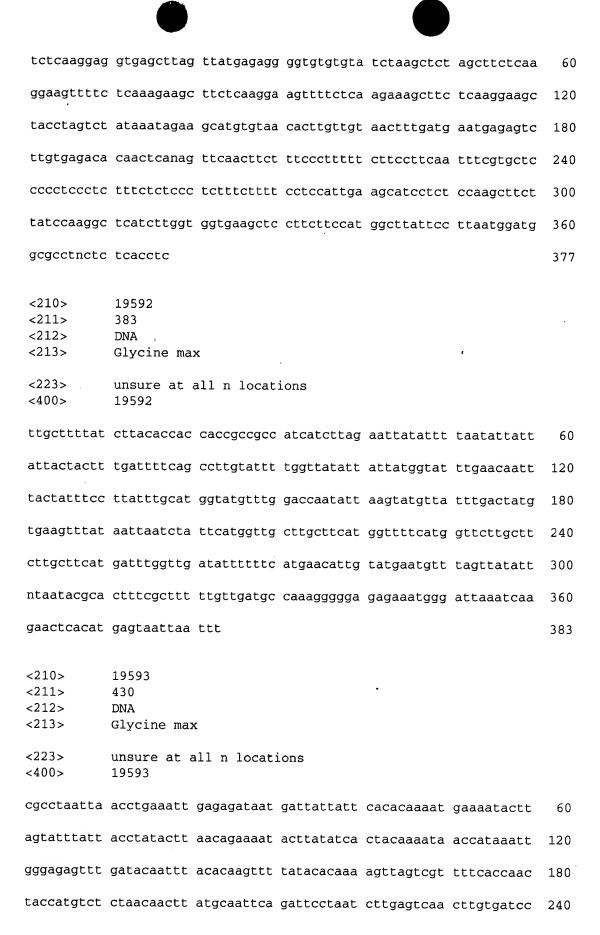
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| ccagacatat | atagetttet egataatett geagattaea taacaaattt aaegtatea | 60 |
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| tatatatata | a tatatatata tatatata tatatata tatatata | a 180 |
| tatatatata | tatatatata tatatattag agactctatt attacacatt atttatttat | 240 |
| ttcttacaat | atgaacaccc cttatattca aataaaacat cttaaaaatc tctctaatto | 300 |
| tgtctcatta | cagagettaa ateeteteat gttaagataa aatateaeta tattteteea | 360 |
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| ttaaatttat | ttgaaacctt ctatattaaa ctagaaatca aattgagtgt aaattcaagt | 180 |
| ttctttaaaa | tggaaaactt atttttatgc attatcaaac ttaaatgtaa acaaatgaca | 240 |
| agaaagaaag | aatgttactc acaaacattt ttatattagc tcattcttta attttggtct | 300 |
| acatccaatc | cttatcttta tctttagtag aatttttact agatgacaca cactaaccaa | 360 |
| cacacacaaa | tatatatata tatatata tatatata tatatata tatatata | 420 |
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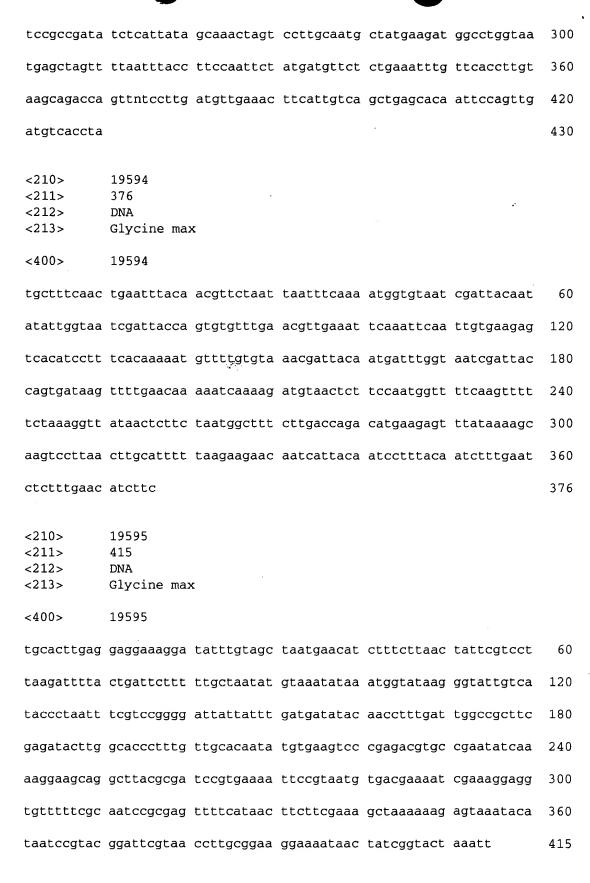
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| aacatctct | a gcacaaatgg tatgacccgt aacggacaaa tctttgtagc gcccgagctt | 180 |
| ctgatgcgaq | g ataaagaccc aaaggggaag gcgaaagtgg gcacgaaata gagcgacaag | 240 |
| gcaagccgta | a ttctggatga ggaggtcccg gccgggaggt ttgctaaggg agaggaagac | 300 |
| ttcggcagaa | aaataatatc cgtagaagaa acaaatgtgt tccttcagat catccagtaa | 360 |
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| ttgccttccc | tagetetaat tttagagett ggaettette atetteetee ggagettega | 240 |
| agttctcctc | attgataact ttcaacttgg agagccaatc taaccctcgc gtatgaactc | 300 |
| ttagccatcc | atgataacca ccgatgacgc cattacggat gcccctgagc tcgttgtctt | 360 |
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| actgtcttt | g atatttggta gttgatattg tgttgtggga ggtaattccg actggatta | a 180 |
| ctcaccato | cc ttcacttgcc aatttgttat gacatttgtt gttggattac ctatgatgt | 240 |
| ttgtttcca | aa gggtagtcta tatcctttct gatggcataa gcatganacc aatcaaagaa | a 300 |
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| gtttgtacc | C ttgtaat | 377 |
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| gtgagagtca | a tttggatcaa tgacaataga tttggatgtt atatgcatga gtatttcaat | 180 |
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| ctcaaggaag | atttctcaaa gaagcttctc aaggaagttt tctcaagaaa gcttctcaag | 120 |
| gaagctacct | agtctataaa tagaagcatg tgtaacactt cgtgtaactt tgatgaatga | 180 |

gagtcttggg agacacaact caaagttcaa attctctacc ttcttcttcc ttcaatgtcg 240

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| cgccccaaa | tgcacaagta | agaagagata | attttccggg | ctctcgtgtc | cgtaaaatgc | 180 |
| attcatatca | tgcatcgcat | aagcatctct | tcataacatc | ataatggaca | tatcctgcat | 240 |
| ttgtccgtta | tcatattcca | gcctcacatt | ttgcatgagt | catggcatca | tcatgcatat | 300 |
| gcgttcaaca | aactttttga | tctgcaaaat | tgcataccat | ttgttttcat | gtttgctcat | 360 |
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| cattcttagt | tt | | | | | 432 |
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| tctcaagaca | attttccatg | atctcaccgt | tgggatcttt | gagaagatat | ctggagtgtg | 180 |
| ctagaagctt | ccgttcccga | gagcatctct | tatttaagca | tttcagcctt | tgctttcgtg · | 240 |
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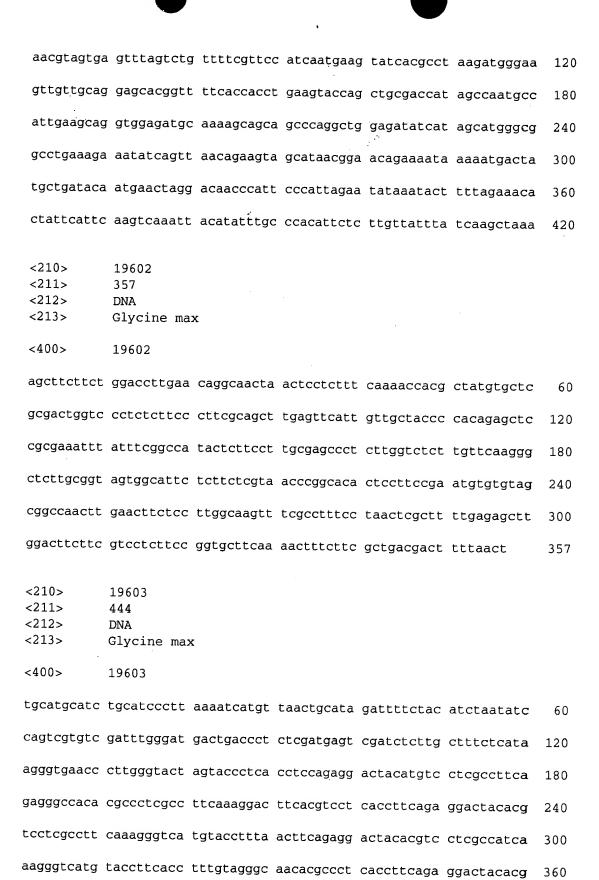




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| tcaacaagaa | tcaagccaag | gctattgtgc | aagcaatcaa | tggggcataa | cacaccaaat | 120 |
| gattatgatg | atggatggct | caaattctca | caaaggtaaa | ctcatcactt | tcaaattgag | ,180 |
| ctttcaaaac | tatcatgaca | tgtagaggag | aatcaaggat | ttcaagtcac | aaaatgtcaa | 240 |
| gaacatttta | ttttcaaaac | aattacccat | ttcttgaaca | tatcctataa | ttcanagaaa | 300 |
| aacatgcaaa | gtcgtacatg | cacacagaat | tgacccaaaa | tattaaacta | gaaatccgac | 360 |
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| gttagcatgg | acctctccaa | ggcaccactg | gaatcagctt | catctgttga | tctttttcag | 120 |
| ttaccagcag | caccatctca | agctccaaca | ttggatttgt | ttcaatcatc | tctttcatcg | 180 |
| gcagatccat | ctttcaacga | gaatcaactt | agtcaaacat | cccatcttgc | atctattgat | 240 |
| tttttttccg | atttttctcc | gcagccttct | actgtaacct | cagatgggaa | ggcactggaa | 300 |
| ttatctgtcc | ctaaaaatga | aggatgggca | acttttgata | tgcctcagag | aacctcctct | 360 |
| actgcacaag | tggaaattcc | aaccactgta | ccctcaaatg | ctaaatcttt | a | 411 |
| <210> <211> <212> <213> | 19598 247 DNA Glycine max | ς. | | | · | |
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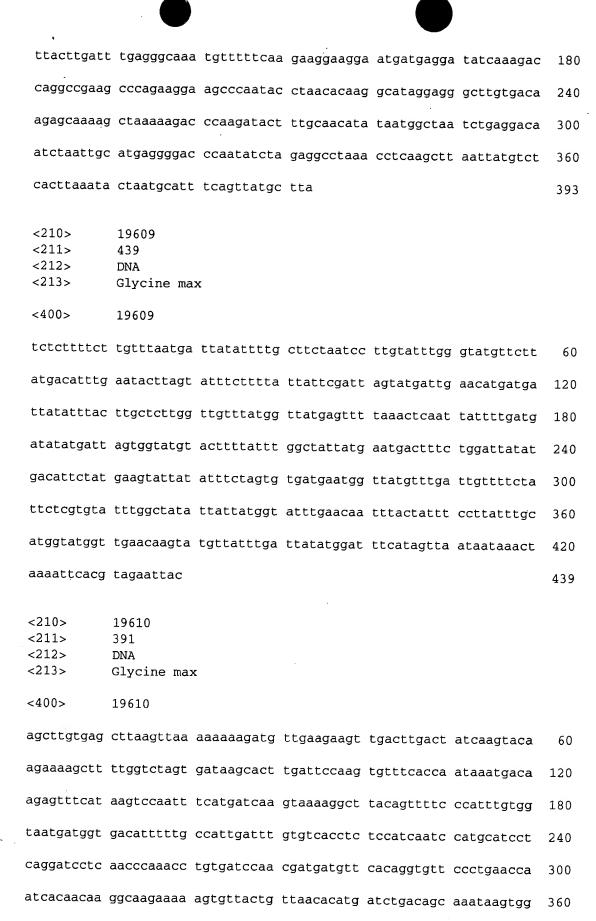
| gctaccacac | c agageteage gaaataaaet atageeatae tgteeeatgg gageeetett | 180 |
|----------------|--|------------------|
| ggactctaga | a tcaagggctc tggcggaaat ggcattcact tatcgtaacc agagacactc | 240 |
| ctttcga | | 247 |
| 04.0 | | |
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| <211> | 153 | |
| <212> <213> | DNA | |
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| tccattatca | a atttctagtg tctcgatata ttacaggtct ctagcggaca tgcaagtgac | 60 |
| atgatattgt | cgcttgaatt tgctcataga ttctcgattc agttgtgagc gtcacgatat | 120 |
| actactggac | acacctcgga catctgatta tat | 153 |
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| <212> | DNA | |
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| | • | |
| <223> | unsure at all n locations | |
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| agttgttgga | Gaagatatot aatototogo ottotototaa kanalaa | 400 |
| agrigitgia | caacatatgt aatgtctggc cttgtgttgg tcaaatagat caacctccct | 120 |
| attaatctcc | tataagagga aacatcttct gctgaaatag gtgaccctga gtgttgatgc | 180 |
| | | |
| ttggtggtgt | aatcacaagg tgtagaaact ggcttagaac caagcatgtc aacattattg | 240 |
| agaatgtgga | gtggstagtt tgtttgstat garatet | |
| agaatgteea | gtgcatactt tctttgatat agatttatac caatagagct tctagctacc | 300 |
| tcaaacccca | gaaagtacct aaagtctcct aagtccttaa ttttgaaagc attgtcaagt | 360 [.] |
| agatntgtaa | ttctttgaat ttta | 384 |
| | | |
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| <211> | 420 | |
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| <223> | unsure at all n locations | |
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| ntgtcctttg | gaaggtgtga tgttntatta gggacttctg aaggtttttg aaagcttgaa | 60 |



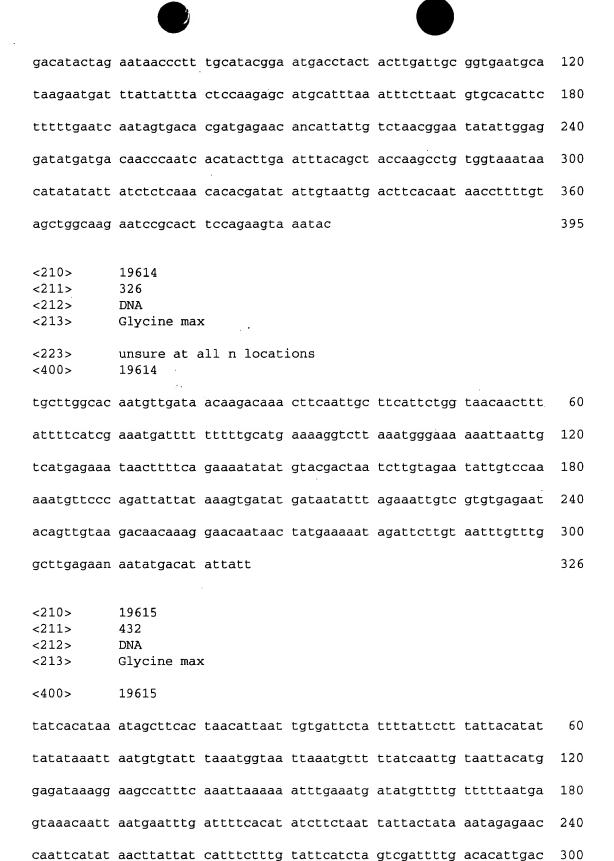
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| gaggactaca | cgtcctcgcc | ttca | | | | 444 |
|----------------------------------|------------------------------------|-------------|------------|------------|------------|-----|
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| <223> <400> | unsure at 19604 | all n locat | ions | | | |
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| ttgattgtag | aatgttaagt | ctttttaata | gaattaactt | gtctgggtta | taggtgtgga | 120 |
| tcttgcaggg | agaagagaga | aagcggtgta | aaacaaaaga | ggcagaggga | agtgaagaag | 180 |
| actttggagt | ttgctgagaa | atggaaataa | aagtggggct | gggtggtcat | ggtcatactt | 240 |
| tagtgtcacg | ttagatagtc | tacatggcac | taaaattacc | aacaatgcac | ctcactaatg | 300 |
| gtgttacttt | aaaatntaac | agaatgacta | ttttacaaaa | cttatgcaaa | gatagagact | 360 |
| attctttaca | tttcanagaa | atagggacta | atgtacaaaa | | | 400 |
| <210> <211> <212> <213> | 19605 445 DNA Glycine max | × | | | | |
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| tccctgcaca | aatttaatta | ttaaatataa | agaaccaatg | cacaagtatg | ctagacaatc | 60 |
| caaataattg | ccaccactta | aacctcccaa | ttattcttca | atttataatt | ataaccaaca | 120 |
| tacagtataa | atatgatgat | taataaattt | ctatgtagta | ttttatgact | taaaaagata | 180 |
| attttaatga | gatattagaa | aaagttctat | tgctaattta | aggtattttt | aaagaactaa | 240 |
| ggagctagga | atcattagta | gtggacccta | agctaagggt | gtattagact | aagtgtagtg | 300 |
| gtgtataagc | aagaataata | taaatacatg | gtaaggcaca | ttgttagcaa | acctaaccga | 360 |
| taagtggtaa | ttatctgccc | tcaaccactc | actcttaaca | ttgtgacaaa | gcattgcatt | 420 |
| taaaccaatt | tctcattaaa | aaaca | | | | 445 |
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| atcgtctact | ttaaagcaaa | tatgcactta | acctttcgtg | acttataaat | aaagatgata | 120 |
| attagacatg | tgagttacac | tttatttaac | ttttcaägat | ttaaatatca | tgtgtcaaac | 180 |
| aaaattcaaa | ttcaaatgat | taatataaat | ggaaaaaaaa | taattgtata | attgattcat | 240 |
| aaatagtata | gattatgttc | aagattaaaa | tatttattgg | acatctctct | ctatatataa | 300 |
| acttaagaat | ccataaacac | gttctgataa | cacatggtga | cacacgataa | tttttcaaag | 360 |
| ttcggatgcc | acgaatactt | aat . | | | | 383 |
| <210> <211> <212> <213> | 19607 435 DNA Glycine max | | | | | |
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| taactaccta | tctcccaaat | gcctttgcga | agatttaata | attaatttgc | atcaatgtta | 60 |
| aattctagat | gttgctaaat | gcgtgggcat | tgagttatca | ttctatgcct | agcaatgcta | 120 |
| acccgatgat | cattttctta | agatgttcta | ttaggtgtta | ccttttccca | agcatataac | 180 |
| ccctaaaact | catgcatgtt | aattcttaaa | tccttactag | gaattaccct | cacccgagcg | 240 |
| aataaaaccc | aaaattaatg | taaggcataa | atgcaagata | agaagaaaag | ttttagaaca | 300 |
| tgatacccta | gaatgaatcc | tctttgcatt | gataactctt | gaagtacacc | atacatcgtt | 360 |
| ggctttttaa | gtttttcagg | ccctagctag | gggattagcc | actcatggcc | attgagggct | 420 |
| ctacaaatgg | gggtg | • | | | | 435 |
| <210> <211> <212> <213> | 19608 393 DNA Glycine max 19608 | | | | | |
| | | gat caat gat | ant good a co | 2224 | | 6.5 |
| | aataggagat | • | | | | 60 |
| Latuutataa | gtcatacttt : | Laarargget | gatctatctc | tttttastat | aggtgatgat | 120 |



| gaaggatcc | a acaacaacca acatacaaca c | 391 |
|----------------------------------|--|-----|
| <210> <211> <212> <213> | 19611 438 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19611 | |
| tctaatgtgt | actcatgtat tgtcaataag tacaatatgt aagaagcaat atgaagaata | 60 |
| aaacatggag | g catggaaact gcattgttgc ctaaacaagt agcaaaaata taacacctca | 120 |
| tctttgagaa | gtgcttactg ttttctcttc ctcagtgacg agaaacttcc cctctgcaaa | 180 |
| atcacaggct | gcaattggta ttgctagctg cttctgcaat ttctttacaa caaacacttc | 240 |
| tgactcttca | catatggtta ccactgtacc attectacct ageegaccag tteggeeage | 300 |
| tcggtgtgca | tagtgaattg aatctgtcgg taagtctaga ttaaccacaa gatcacattc | 360 |
| tgccacatcc | aaaccccttg ctgataattc atttgtaacc agaactctca cctcaccatt | 420 |
| cttgaatntc | ttcagagt | 438 |
| <210> <211> <212> <213> | 19612 195 DNA Glycine max | |
| | | |
| | cgctgtgcgc acgcattctt gttcctgaac gccgtacact gtacgccatg | 60 |
| | tatgtgctgt cagacgagga cactgtttgt aacttgcata tctagagtat | 120 |
| | ctcacacata gcttgtcaaa atttggatat aggatactcg cacttgcaag | 180 |
| cgatatgcgt | ctatt | 195 |
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| agcctgcatg | acttcggcgc tctaggtact tctttacatt gctatatctt aatataatat | 60 |

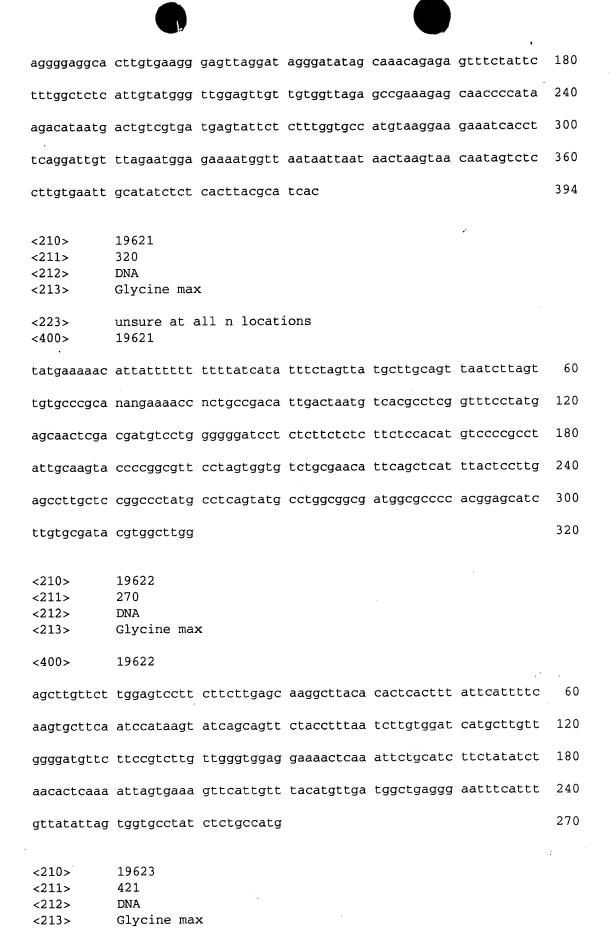


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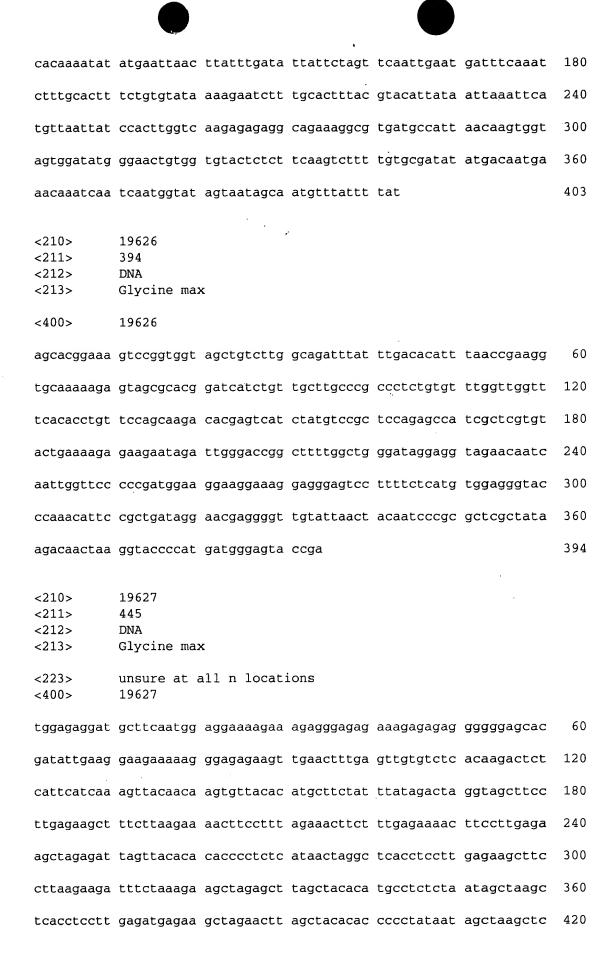
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| actataatca | ta | | | • | | 432 |
|----------------|--------------|------------|------------|------------|------------|-----|
| | • | | | | | |
| <210> | 19616 | | | | | |
| <211> <212> | 395 DNA | | | | | |
| <213> | Glycine ma | x | | | | |
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| tggaggaatc | ttctggaggg | cccaagtggg | cctggttgct | atttacaccc | cccgtttact | 120 |
| aaatgcaccc | cctttctatt | tttttgtaat | tctttttccg | taatgttacg | aaactttacg | 180 |
| aatttcgtaa | cgatacctat | tttccttccg | caaggttacg | aatccttacg | gattatttat | 240 |
| ttactctttt | ttagctttcg | aagaagttac | gaaaactcac | ggattgcgca | aaaacacgtc | 300 |
| ttttcgattt | ccgccacatt | acggaatctc | acgaatcacg | caagcctgct | tcctttcaat | 360 |
| ttctgagacg | tctcaagact | taatttattg | cacgt | | | 395 |
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| <211> | 428 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | ζ | | | | |
| <400> | 19617 | | · | | | |
| tgatcaaaac | aattatctaa | tcattcctat | ccactcaatt | catacaattg | ctcattcaaa | 60 |
| taattatcaa | acactcattt | cataccaaac | aatccattgc | atatcatttt | caatcaattc | 120 |
| attgttcaaa | cacgettttg | gtacaagcaa | acaactcaaa | gtgctgaaat | ttaaaataac | 180 |
| taaaatataa | agcaaactaa | atactaataa | actaaaatgt | tcatgctttg | cagaaattaa | 240 |
| actaaacaca | atttaaacat | cctgctcatc | ttgtggctga | tcttcattat | gatctagtgt | 300 |
| tggagctgct | gatgaatcct | ggataggctg | ctctggctcc | gtgactggtg | tagatggctg | 360 |
| ggtctcctca | agaactggtg | caagagatgg | cttaagtatt | tgatctatgg | aagtcccctc | 420 |
| ctcctgag | | | | | | 428 |
| 24.0 | | | | | | |
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| -212 | Claraina mar | | | | | |

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| ggtgattttc | aaccatggag | ttgcagtgga | agataaagga | gaaatgggga | gaggaggcgc | 120 |
| catacactag | ggaataagcc | ttggaagatg | aaggttcacc | accaagagag | tgtcttggat | 180 |
| aagaatctta | gagaggaagc | ttcaatggag | gaagagaatg | agaaagagag | agagagagag | 240 |
| agagagagag | aaagtggcat | gtaaaattga | agaaagaaag | gtagagaatt | tgaactttga | 300 |
| agtgtgtctt | acaagattct | cattcatcag | agttgtgaca | agtgttacac | atgtttctat | 360 |
| ttatagccta | gccaatgact | aaatgaaatt | ttatttt | , | | 397 |
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| attggttgtt | atcttggttg | ttgcatctta | gtacatttga | tattţgtttt | gcattgtgca | 120 |
| tcatcatagt | gtgtgtgaag | aaaagtttct | aagttataaa | aattacttta | gaggcaaaaa | 180 |
| ctctttattt | taattgatta | caacctcatt | gtaattgatt | acaacaagct | attaaagctt | 240 |
| gtagagttaa | gtatcgtatc | ggtttaatcg | attaccgata | tctcataatc | tattacacta | 300 |
| ttgtttgaga | caatgactga | tttatttagg | agtctttgct | ttaatcgatt | accaagtgga | 360 |
| ttaatcaatt | acttctatct | cgttcaagtg | ttctggggtg | aacaagaaca | ctttaatcaa | 420 |
| ttactt | | | | | | 426 |
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| agagagetet | gcaaccaaat | cctttatage | naacatnnaa | aaacaatato | ttataaqqaa | 120 |



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|-------------------------------|--|-------------------|------------|------------|------------|-----|
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| gggcccaagt | gggcctggtt | gctatttgca | ccccatttt | tactaagtac | accccctgc | 120 |
| ctttttttg | gtgattcttt | ttcgcaaagt | tatggaaact | tacgaatttc | gtaacgatac | 180 |
| ttgttttctt | tccgtaatgt | tacggaacct | tgcggattac | ataatcatcc | cctttttgac | 240 |
| ttacggaatg | ttacggaacc | tcactaattg | tgcaacgatg | cttccatttg | atttccggtg | 300 |
| tgtcacggaa | ccttacggat | tgtgcatcaa | tattttcttt | tgttttccgg | catgtcccgg | 360 |
| aatttcacaa | attgcctaat | gatgggttcc | aagcacctca | caaggaccaa | acaaaagttg | 420 |
| С | | | | | | 421 |
| <210> <211> <212> <213> | 19624 395 DNA Glycine max | ĸ | | | | |
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| taaattgtat | ttaaaaaata | tgtttataaa | tttaattttt | tctaaaatat | taaaataata | 120 |
| aaaaatttat | tatattttt | agatattaat | tttttataa | ttaaatttac | aataagaaaa | 180 |
| aaattagata | tatttttcag | atataattct | taataattaa | ttttatgata | attatttgag | 240 |
| taatttaata | atatgaaaat | ttatttattt | taaacaatat | attataatta | attaatttaa | 300 |
| attgagtagt | attgcggaaa | ttcttttgtt | attagtcaat | cataagtaat | cctacaatta | 360 |
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| | tctaattatt | ctcaatcatt | tataa | | | 395 |
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| acccccatg | a caaanaacat gaaaa | 445 |
|-------------------------------|--|-------|
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| <400> | 19628 | |
| agcttgtact | t tgatcttgga ttaatgagct gaatcatagc taacattaac taatcataa | = 60 |
| tagagaaact | t ttcgctccaa aatttgcctc caaaaattca atttcaaatt caagtgaaat | 120 |
| ttgaatacaa | a attcagattt ccctccaatt ttgtgtgaca cttacgctat aaatagacgo | 180 |
| catgcgcgcg | g catatgtteg actgegatea tttgaaaatt acaetteaaa tttetgaeet | 240 |
| tattttaago | acteattgeg egtegtteta ttetetgeet tatteaactt ettecaeate | 300 |
| taccc | | 305 |
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| aagctttaga | attataacat aagaactgtg attattgaag aatctatcca tgttgttttt | 60 |
| gatgaaattg | accctatatg gccaagaaag gatacacttg atgatattgc tgatacatta | 120 |
| gacgacatac | acattgatga gaaagggcat agaggcaaag gaaatggtaa tgaataagac | . 180 |
| tgtcatattg | atgaaaataa aaaaaataaa tatagatett eeaacagagt ggagaaette | 240 |
| aagatatcat | gctcttgata atatcattgg tgacatctca taaggggtaa caacttgaca | 300 |
| ctctctcaaa | gatgcgtgcg ataatatgac tttggattcc ttaattgaac ctaaaaattt | 360 |
| atatgaagcc | ataattaatg aacactggat tattgctatg caagatcagt tatatcaatt | 420 |
| tgaaagaaat | aaagtctggg aa | 442 |
| | 19630 385 DNA Glycine max | |
| <400> | 19630 | |

| agcttttta | t taattacact | catactgtaa | a togattacta | a gaggagattg | tcagaaaata | 60 |
|-------------------------------|------------------------------------|--------------|--------------|--------------|--------------|-----|
| ttctcaaca | g tcacatattt | tcagttggtt | cttgaatgg | c catcaaaggo | : ctatatatat | 120 |
| gtgacttgag | g acacgaattt | gaaaaaaaga | gttttcataa | a caaaaaggta | ttatcttctt | 180 |
| aaaaagcaaa | a atcattttat | cctctttcaa | gagagatato | ttettetett | cttctttatt | 240 |
| aggaaaaggg | g attaatagac | : tgatggtctc | : ttgttgccaa | a gaaatctgaa | cacataggaa | 300 |
| gggttggcct | tgtgtggtgg | agatcttgta | gcaggctgtc | c acaatatagt | ggaactctca | 360 |
| atcaagttgt | ttggggactg | gacgt | | | | 385 |
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| tggaagttac | accaagtgtt | gcacatgctc | ctatgaccag | ctaggttgct | tgcttgtcta | 120 |
| actgtcttga | catcacttcc | ttgataagct | tctttaacaa | aacttccttg | agacactcga | 180 |
| gcttatctac | tcacacgcct | ctaataacta | cactcacctt | cttgcgaagc | ttccttgtta | 240 |
| tgattcatag | tgaagcta | | | | | 258 |
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| tgtaagtgct | atgcagtccc | aagagcagcc | gaggttgatg | agacaagacg | gtgtagttga | 120 |
| aggtgcttta | ccaaccaagc | accatcatgc | tgaatccagt | agaaagaaat | atgtcaagaa | 180 |
| gaaccagcaa | acaagcagcg | aaaattgtgc | aaacaaccaa | aacaaaggta | agggtaaaaa | 240 |
| gaaaaattat | ccaacttgcc | agcattgcgg | aaaattgggt | cacccaccat | acaaatgttg | 300 |
| gaaacgacca | gatgcaaagt | gctgcaagtg | caatcagctt | ggacacaaag | ctataatttg | 360 |
| tagaagcaaa | tttcagcagc | atgaagtcga | tgcccaagtt | gttg | | 404 |

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|----------------------------------|------------------------------------|-------------------|---------------|------------|-----|
| <223> <400> | unsure at all r | n locations | | | |
| ctcaagctto | ttagggtttc aagc | caatggt actagggt | it caagetgtgg | tgatagggtt | 60 |
| tttgaggaad | cgtccaagaa gagg | gatgate ttttaegte | ca aagcaatagg | aaggtcaaaa | 120 |
| taaaacctga | cgaggatagc acca | ctctcg atatagagg | yt tgagaaagaa | aaaggacgtt | 180 |
| gaagaagagg | tggatcccta caaa | aagaag ttgttgtca | aa ataacattgc | agtgcctttc | 240 |
| tctccttttg | atggaagatg tttt | gataag cctaacaad | a aaatccccc | tgccaccttt | 300 |
| gtcgaaagaa | gatagggaaa gatt | ttacga ggttaagat | t acaaatgagg | agttcagtga | 360 |
| ttggtgtaaa | ccctggaagg gctc | ccttat ggtgatatt | g ttgggtaaga | | 420 |
| ttgtatgatg | gagacaaagt tgaa | ta | | | 446 |
| <210> <211> <212> <213> | 19634 384 DNA Glycine max | | | | |
| <400> | 19634 | | | | |
| agcttaagtt | atatcataca tatta | agtact ccttttata | t tctaccatat | ctgatagctc | 60 |
| actttgtaag | aggaaatgag aacat | ttttct cgaatcaca | a catagatcat | tcatttatta | 120 |
| | tgaagcgact agaga | | | • | 180 |
| | tgaattttag tgaaq | | | | 240 |
| | ctgtctcagg acgta | | | | 300 |
| | gtcagattga agatt | taaagg ccactctca | a cataatacta | agctċaaacc | 360 |
| tccaatttct | ccatcattga agca | | | | 384 |
| <210> <211> <212> <213> | 19635 438 DNA Glycine max | | | | |

| <400> | 19635 | |
|-------------------------------|--|-------|
| cgctatgaa | c atcaggcatg atcttaccaa ttaattagtt gattgagtgg ctagtatgtg | g 60 |
| tcggtgttg | c tgtcatccaa ggtatcatca ațaattagat cgaatttgct tttataaaaa | a 120 |
| attactgca | g ccaacaatat ccacgataaa aagaatagaa tggcctacaa tagtctcata | 180 |
| catttatca | a gctacagcat ccagactaac ggttgcaatt tgcccttctt tatacaagtt | 240 |
| attattacco | c aaggacaatc tataacctca tcccacacat agcagtcgta tagtttgagc | : 300 |
| caaataggaa | a tttttagacg agacatggaa agggaatata catgttcacc aagcataagg | 360 |
| atagtggtag | g agctgggata aaatttctgc tgttatttat taacaatccc tacaaagggg | 420 |
| tgatacttat | aaacaaag | 438 |
| <210><211><211><212><213> | 19636 250 DNA Glycine max | |
| <400> | 19636 | |
| cgctttcttt | tatatgtgca attaatagaa aggtateeta tagtagtegt ggaegtaege | 60 |
| acatacattg | tgtgtcgatc cacattaaaa catgcgactc tcctctcttc cctttattca | 120 |
| tacacctgct | tttattgcac tatactaaca tgcctcacca tgacaacatt ggccaaccaa | 180 |
| gtggagtatc | tgacttctat gatgaactgg actatgagga gcttgacaat ctctactcat | 240 |
| atcgctttac | | 250 |
| <210> <211> <212> <213> <223> | 19637 422 DNA Glycine max unsure at all n locations | |
| <400> | 19637 | |
| | aggtccagga aagatattgc ggncgaattt actagttccg cccgtgagta | 60 |
| | cgctttaaga gcgctgtaca ccagcatcgc ttcgaggcca tcaagggatg | 120 |
| | cgggagcgac gcgtccagct cagggacgat gagtatactg atttccagga | 180 |
| ggagataagg | cgctggcggt ggacatctct ggttaccccc atggccaagt tcgatccaga | 240 |

aatagteett gagttttatg ecaatgettg gecaacagag gagggegtge gtgacatgag 300

| | gtcctgggta | aggggtcagt g | gatcccgtt | tgatgccgat | gctatcggcc | acctcctgcg | 360 |
|---|----------------|---------------|-------------|------------|------------|------------|-----|
| | atatcctgtg | gtgttggaaa a | atgccagga | ttgcgagtat | tggtcaacga | ggaaccggtc | 420 |
| | tg | | | | | | 422 |
| | | 10.500 | | | | | |
| | <210> <211> | 19638 329 | | | | | |
| | <212> | DNA | | | | | |
| | <213> | Glycine max | | | | | |
| | <400> | 19638 | | | | | |
| | ttactttcta | gcactaacaa g | actgagtat | ctcaagacca | tcatcttgga | gatactccaa | 60 |
| | gctccaacca | tagacgccag g | aaggtcatg | gtgggagaaa | tggatgaacc | tgactatatg | 120 |
| | accccctata | agaatttctt a | atttgaggg | gtgtttccac | caaacaagaa | tgaaacctga | 180 |
| | cgccttaaaa | ggaaggctag c | ttctatgtc | atcattgact | gtgaactatt | gaaaagagga | 240 |
| | ttaacaacac | ccttgctcaa a | tgccatata | gccaacaagt | agactacatc | atgcgagagc | 300 |
| | tacacgaagg | aatttacagc c | tccatatc | | | | 329 |
| | <210> | 19639 | | | | | |
| | <211> | 433 | | | | | |
| | <212> | DNA | | • | | | |
| ٠ | <213> | Glycine max | | | | | |
| | <223> <400> | unsure at all | l n locati | ons | | | |
| | 11007 | 15055 | | | | | |
| 1 | ntggtctcta | caaatcttcg ca | acagcataa | tctatcaatt | tctctggaac | ttggaccttt | 60 |
| (| ctctctctag | aaaaacctca ca | atgcagaag | ctccttgaga | aaaatggcca | aactcgcttt | 120 |
| (| ccaaaatctg | atttcaggct ta | aataggtg | gctttgttca | tgcttgtgtg | cttagcgcaa | 180 |
| t | tctgaatcg | cttagcacgg ag | geggatgga | ccgaagcggt | gcgcttagcg | ggatggccct | 240 |
| t | cactcaaca | aacaagcaca ac | ctcatcctt | cttccagatt | cttccttgcg | cttagttgag | 300 |
| ç | gaatgttgcg | ctcagcggat ag | rctcactaa q | gccagcagat | tggcttatcg | agagggtgaa | 360 |
| a | atcaacact | tcaaaacttg cc | taattatc (| ctgaaattga | gagaaaatga | ttattaaata | 420 |
| C | acaaaatgg | gag | | | | | 433 |
| < | 210> | 19640 | | | | | |

| <211> <212> <213> | 381 DNA Glycine ma: | x | | | | |
|-------------------------------|------------------------------------|------------|------------|------------|------------|-----|
| <400> | 19640 | | | | | |
| tgtgaaaccc | accatgcgtg | catagaaagg | tcgtggtcgg | agcgcgaaac | tgatgcccct | 60 |
| ccacagtgtt | atgtaagggt | ataccaccaa | ctacttgcca | ctcaaaggca | catccgaggc | 120 |
| aaagcctgaa | gtcgcttaaa | ccatggcggg | gaatatcatg | aagatacccc | aaggattgag | 180 |
| agacaagtac | atgacgagcg | agacggctat | caaagaggat | gggagcagaa | acattgacat | 240 |
| gctcactggg | agtgtacgcc | ccactagaca | ggagtctaaa | ggggaagcca | accatatggc | 300 |
| gggaatgcgc | gcgacatctg | aatggcacac | agaaggcctg | tcggagcacc | aaatcctgcc | 360 |
| tacatcaatg | ccgaggatgt | С | | | | 381 |
| <210> <211> <212> <213> <400> | 19641 312 DNA Glycine ma: | × | | | | |
| tcgctgcggt | attctatatc | gagcgtacgc | ttatattacg | agttttattc | cgacatacga | 60 |
| ctataatgtg | attgtcttct | gcatttgctc | ataacttcgg | tgtacaattc | cgagtgtcga | 120 |
| cgacatacaa | cgggactcaa | tccgacgtcc | agatcaaaag | ttgacgtcga | ttgaattggc | 180 |
| tatcagcttc | ggtattcatt | ttcgagcgac | tacatagatt | acgttacctc | ttccgacatg | 240 |
| cgagtcgtca | cgaattgccg | agcgactttg | ctcatagcgc | cagctttgta | tttctagcat | 300 |
| atggagatat | ta | | | | | 312 |
| <210> <211> <212> <213> | 19642 364 DNA Glycine max | ĸ | | | | |
| <400> | 19642 | | | | | |
| | | gacaataagt | | | | 60 |
| | | ttgaatgtcg | | | | 120 |
| ttttactcac | atgtctgatt | gagtcccgta | acatatagag | acgctcgaaa | ttgaatgttg | 180 |

| aagctctga | g ccaattcta | a cgaccatato | cttttactct | ggtatctga | t taagtccggt | 240 |
|----------------------------------|------------------------------------|--------------|--------------|------------|--------------|-------|
| aacatatcg | t gacgcatga: | a attgaatgtt | gaaaccctca | ctcaattca | a acgataataa | 300 |
| ctatttact | c tgatgtctga | a ttgagtcccg | r tagtaacacc | agacgctcat | aatgaatgat | 360 |
| gaac | | | | | • | 364 |
| <210> <211> <212> <213> | 19643 381 DNA Glycine ma | 1 x | · | | | |
| <400> | 19643 | | | | , | |
| cactatacaa | ı tactcaacgo | tgcacatgga | acttcgagcg | cacctaatgt | gacaggattt | 60 |
| ttcagacato | : cgagtaaaaa | gggattgtca | tttgaatttg | ctcagagcat | caacattcaa | 120 |
| ttacgagcgt | ctagatatat | tacgggactc | aatcagacat | acgaataaaa | agttattgtc | 180 |
| gtttgagttg | catacagggt | caacattcaa | tttcgagcga | ttcgttattt | tacgggactc | 240 |
| aatcatacat | ccgatgaaaa | tggatatgtc | tttgaattgg | ctgatagctg | caacattaaa | . 300 |
| ttttgagcgt | ctcgatatat | gatgggactc | attcatacat | ccgagcgaca | atgttttgtg | 360 |
| tttcgaattc | gttgagagct | C | | | | 381 |
| <210> <211> <212> <213> | 19644 351 DNA Glycine max | · × | | | | |
| <400> | 19644 | | | | | |
| tagcgcatta | catctaacaa | gttctatgtg | tattctttgg | tgatgaactc | tccttgccaa | 60 |
| ttttgctact | atttactaaa | cttaatattt | gttagctgat | tccttcttaa | tctaggtatc | 120 |
| gatcatcgac | cagttgactc | actaagtaac | ccactatttg | attcattgtc | ccctacatat | 180 |
| ctaactaacg | aacctatcaa | acaacctctt | aagttgcagc | cccaggggca | gtgctgctga | 240 |
| ttcattgccg | gaggggcatc | aattgagaga | tttgggtggt | attgagggtg | cagccacagg | 300 |
| gatagagcct | actgattctt | ggtcaaccat | aatgggcgtg | aatatgcttg | g | 351 |
| <210> <211> <212> | 19645 411 DNA | | | | | |

| <213> | Glycine max | • |
|----------------------------------|--|------|
| <223> <400> | unsure at all n locations 19645 | |
| tgccccncag | g ctcgcccagg agagcaagtg tggcttcctc tataagcaac cgccttctgg | r 60 |
| aggaagaato | tgaaggccca agtgggcctg attgctattt gcacccccat ttttactaaa | 120 |
| tacaccccct | tgctcttttt tggcgattct ttttccgtaa cgctacgaaa ctttatgaat | 180 |
| ttcgtaacga | tgcttgtttt ctttccgtaa tgttatgaaa ccttacggat tacgtaatca | 240 |
| tcccttcttt | gccttccgga atgttacgga actttgcgga ttacgcacta acacttcctt | 300 |
| ttaatttccg | gcatgtcaca gaacttcgcg gattgcgcta caatgctttc ttttgactcc | 360 |
| cgacatgtca | tggaacttca tgaatngcct aacgatgggt gccaagtacc t | 411 |
| <210> <211> <212> <213> | 19646 353 DNA Glycine max | |
| <400> | 19646 | |
| acatgtatgg | acggtataaa tagtgagatc taatcaattt actagatcaa taagattaat | 60 |
| actacatata | ttttaaatca aagaaactaa caatgtaatt aaattcaaac ttagttcaaa | 120 |
| agtaaatatg | cgtaaatgtt ataatttctt ttatgttaat tcactttgaa aaatcagaat | 180 |
| taatttttt | aaaaataaat aatattaata atgaatttta atttacagct ctgcaataca | 240 |
| tactttcatt | gttggaacat caatatatat ttgaataatc caaattacag aatatgtgaa | 300 |
| gaatatatga | gatatctagt gaaacatgat aattcatgaa tatatcttaa tat | 353 |
| <211> <212> <213> | 19647 418 DNA Glycine max 19647 | |
| tttcaaacgg (| gtaatatget cacattetet ttettetata teatatteaa aettgteeaa | 60 |
| | aagtcatatc gactcacaga aagtcatatc agtctcatac aattaatata | 120 |
| | cctaatgtca catcctatca gagcgtggtg cttccgtgtc ctctagcatg | 180 |
| | agtcatcca cctattcatc tgctccccg aacacaaggt caagatcatc | 240 |

| acaggatcca | a aacacaacaa cacacaggga gtgagttatc acattcctag ctaatagata | 300 |
|----------------------------------|--|-----|
| aacatgacaa | a ttaaatatto gtattatata aatgagatao cacttgotta aacatacoto | 360 |
| acgtaactto | accactgtct cattcaaaat gcacttctca ttataatcac attacaca | 418 |
| <210> <211> <212> <213> | 19648 401 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19648 | |
| agctttataa | gcgcgggtct gggagacaaa ggtcaagcgt tcgcgatatg cgaggatgat | 60 |
| attccgagta | ctttggattt ggtacgacca tgccctcctg atttccagct gggaaattgg | 120 |
| cgagtggagg | aacgccccgg catttacgca acgagcataa tgtaaacctt tacggtttta | 180 |
| adagctctat | agttgggcct aggctttaga gtttttcctt ttgttaaggc tttgtgtctt | 240 |
| ttgtttttga | atttataata caaggatett tetteatetg tteetggtet etacceatte | 300 |
| tcattcattt | gcatgtttac ttctttntct gaaacggcag atccggtgac gagtcccccg | 360 |
| aaggtactta | tacctgngac ccgcctatcg acttcgagcg a | 401 |
| <210> <211> <212> <213> | 19649 445 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19649 | |
| ttggctacaa | ccttttctcc ccctttggca acatcaaaaa gccaaagaac tcggaaatca | 60 |
| acacagttat | aacaatggag tagcaagata taagtatcag agtattaaat ccaataagcc | 120 |
| aaactcataa | tcaagaaaat aatcaaacca gaattcaaat aacataaaat gtcaacaatc | 180 |
| acaaaatatc | caagaccgaa acacaagaaa aataagcaaa gtacttagca taataatgta | 240 |
| aattctaaga | aactaaaagc caaaatacac ggcttataaa agataaataa gcagaatcta | 300 |
| aaatctaaga | agacggagga ggtggtggaa gatcaaaact ctgacgaatg tatccgacat | 360 |
| cctcttcaag | ctgtgtaaga cgaatgtcca taccggcaaa gcgtgaatct aacgagtcan | 420 |
| agcggtcacc | aacatacgaa cgaag | 445 |

| <210> <211> <212> <213> | 19650 384 DNA Glycine ma | x | | | | |
|----------------------------------|--|------------|-------------|--------------|--------------|-----|
| <400> | 19650 | | | | | |
| agcttattca | catgaaacac | aaggcatat | t tatgtetet | g gtgagcctaa | a gctcttcgct | 60 |
| ctagaggaca | cgaagctgtg | caaggaaag | c agcacatag | a tccaagaaaa | a gaaactgtgg | 120 |
| accetgeace | ctttgaagat | caagcagata | a cttttcttc | c cgggttttgt | aaagctgttg | 180 |
| atgtgactgt | aagaattaga | gaatattcto | gaaaagaati | t ataactacco | : aataataaca | 240 |
| gaagaattag | tttattgaga | caatctgtct | tataattggt | tataacactg | taaagaaaca | 300 |
| gcaatgattt | acacgtgtat | actgtctagt | attataaaca | ı tggagggagg | gaagcaaaat | 360 |
| aatcagaaga | ctatttagtt | atta | | | | 384 |
| <211> <212> <213> <223> | 19651 429 DNA Glycine max unsure at a 19651 | | ions | | | |
| gacttcaccc a | attntagcat | attcttaccc | aagttgtcca | agcaagagct | ctacgcctcc | 60 |
| caaaaaaaag (| ctccaaattt | tgtttagtac | taatatatct. | caaaaaaatt | attgtaaaat | 120 |
| tatatttaaa a | ataaatttt a | aattaaaata | ttataagtaa | ctattaatct | ttttattaat | 180 |
| aatgtagata a | | | | | | 240 |
| aaagagattt a | | | | | | 300 |
| aaaaaatatc a | | | | | | 360 |
| aacaaaatct t | caaaaatca a | acacgattg | tatcaagggc | gttaagataa | aacatgaaat | 420 |
| tgttctttt | | | | · | | 429 |
| <211> 3. <212> DI | 9652 52 NA Lycine max | | | | | |

| <223> <400> | unsure at all n locations 19652 | |
|----------------|--|-----|
| agcttttato | c tttattgcac tattgaccta cctaggagat tttataccta gaggtactca | 60 |
| tgattnatgt | cgacaagtac tacatatttg caaaaccaac tgatgagtcc tcctatgact | 120 |
| aggcatatct | ccccatacga agtattttgc ccgaatacga gtatttaaat taactaaaat | 180 |
| aaaccaatta | aaaatacact gtgcaagttt gattcataaa aatactaatg aaatgtaaat | 240 |
| gactctcacc | attttaaatt aatgaaatat ataataatat taattaaggg aaatatgatc | 300 |
| aatataatat | ttgtcggtta aacaatcaaa ttgataacac aaggatagct ta | 352 |
| <210> <211> | 19653 415 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <400> | 19653 | |
| catactctat | cagctgatat gctcagtgta ggtttactat cttgaacaca tttcattcaa | 60 |
| | taaagcatag gtatgacttc caacataaca agatacatta ttacttgggt | 120 |
| | ttaacatgat tactttaaaa tattaaatga cagacatcat atattgaact | 180 |
| | gtatcctact atatgtaaga cgggtttacc ctagctgata gagcccaact | 240 |
| | gatttagttt acgaatctgt tatcaatcct ctctttagga cttatctttt | 300 |
| | tttaaggagt tggttgaggg agaggttaat aaatatatga cagaatgtta | 360 |
| gtgaggaaca | ccttctggtg aatttctgat gcatttatat tttagtaata atact | 415 |
| <211> <212> | 19654 388 DNA Glycine max | |
| | unsure at all n locations 19654 | |
| agcttgtcgg t | gtttgaaat ttcttctttg gtgttgaaga agtaattatt agaggcttag | 60 |
| tgaagttgga a | atgttctaac tagcttgttg tgcaattttt gccatgccct aacaaagtaa | 120 |
| | atgcgttta ggaacccatt aagataatgc tttcgtgttt atgctcccta | 180 |
| gtgttctttg c | ecttttcttc tttggtgttg aagaaattag tattaagggt ttagtgaagt | 240 |

| tgaaatgtto | c taacttgttg tggaattatt gactttccct aagaatcaat ttgacttgc | g 300 |
|----------------------------------|---|-------|
| cttagctttg | tggaaacttc aatcagtaga tagattcaaa tctcttgttt gaagtattt | c 360 |
| ttcagcctgt | ngcatgtgtt acttggag | 388 |
| <210> <211> <212> <213> | 19655 447 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19655 | |
| ctcaagcttg | gaggtatcaa aaagattatc ttgttataag gtctaattct cctagctaga | ı 60 |
| | caacattgga ggaggtggca ggtagagagg gagagtggga cgttggaggg | |
| | acaaaaagaa gaattatcaa ttgaacatag gtcattatat tggaaaacaa | |
| | actttttctt cttgcatttt ccagtatatc atacataaaa aaacttatgg | |
| | tgaagtagga gaaagtgtgg gatatctata tctcaccaag gaagaacaag | |
| | aatacgattt tgtttggttc ttggaggtga ctaatcccaa aaggcttgaa | 360 |
| tatcaacttg | ataacattca tataagagac ataaagttgt ttgtttactt aactagantt | 420 |
| gggatgacaa | caacaaagga ggtgaag | 447 |
| <211> (212>) | 19656 394 DNA Glycine max | |
| <400> | 19656 | |
| agcttgagcc g | gttgttgatg agtactttga ccataacgtg ttccatgcat ttgactgaca | 60 |
| catgtaaagc c | ectgttatgt cetetteeet egaeggggat tteetettea geaaatgtga | 120 |
| ggtagttgtt g | gctgtgatg ttattgacga tcccccccaa agccttctac ggagatgtct | 180 |
| tggactacgt g | agcttcatt caagaccttt accagcaaag ctcgatgagg ctcagagctc | 240 |
| atgagcagct c | cagaaggga gactctagct agggtcttgt tgagttgttc aatgactttg | 300 |
| aactcgcttt g | ttggataat gcggaggaac tccctcctcg agtgatacct ctttcttgcc | 360 |
| acagctctct c | ttcccatgg aaagatcctt cgtc | 394 |

| <210> <211> <212> <213> | 19657 419 DNA Glycine max | |
|----------------------------------|--|-----|
| <223> <400> | unsure at all n locations 19657 | |
| cactcagct | t gtaaaaatgg aagaaaagaa accgaaggtg aacgaatttt agatgaatgt | 60 |
| ctaaacaac | a agaaatgaat tgaaagtete ggatteaaaa aettaeeegt tgaagaaega | 120 |
| agaacgaat | g aagaacggat gaagaatggt gaagaacgac ggaaaacctt cacggatttg | 180 |
| ctcacggaa | a agtettggaa gtgttaegga aacaeetegg ettggatttt etteaeggaa | 240 |
| acaattatt | t tcaccaaaaa caactgaaat gtatagnnaa ggaggttaga gatatttgga | 300 |
| acageetee | c ttcgccaatt tataggaaaa ggggggagga cgttgtcgtc cagtgtgcct | 360 |
| tgaaaattt | g aacaccgcta teegeaceee etetegataa gtteacgntt ttetttegt | 419 |
| <210> <211> <212> <213> | 19658 383 DNA Glycine max | |
| <400> | 19658 | |
| tatcttgtta | a aatcagggaa acatgctgaa ctctggggtg ccctccaagg aaacatactc | 60 |
| atccagatca | acatggcgaa ccagtttaag aacatgctga ttgacaataa tggcaaaggt | 120 |
| gggggaaaca | acaacggtca aaagggtggt ggtgggaata accagccaaa gggtaataat | 180 |
| caacaagggc | agaaccccca acagcaactt catcagtatc tgcaacagct tcagcatatg | 240 |
| aaagggttcc | aagatetgaa getgteteaa tteaatgaea tgaaaetgee caaceegaae | 300 |
| ccgaacccga | accegatagt eggtaaatta aatttgeetg aegaggatga tttgtetgat | 360 |
| gatgaaatat | actagtttga tga | 383 |
| | 19659 402 DNA Glycine max 19659 | |
| ccttctcagc | ccacgcattc atagtatatt gatcattaat attcacacaa ataacagaat | 60 |
| | | |

| • | | | | | | |
|-------------------------------|-----------------------------------|--------------|--------------|-------------|--------------|-----|
| ctatcccct | t agccttaaa | c ttatcaata | t tctccttgt | a aggaggaac | a tgtttgtttg | 120 |
| aacaaactc | c tgtgtatgc | a ccctgagag | a aagccaaag | c aaaaaatac | a ttactatgga | 180 |
| ataagtcta | t aacaaaaa | a aagtaacac | a tttctttca | a acaattcca | t ttcctataco | 240 |
| ataagaagc | a ttcattcaa | c cacataaaa | t agtaccaaa | t tttacaaaa | c aagaccaaca | 300 |
| aacaccaaa | g agacacccc | a aaacatatc | a acatatcaa | a aaaaattgc | a agttaaactt | 360 |
| actgggagc | c caaagatga | c aactttctt | g teetgaaac | a tg | | 402 |
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| agcttttagc | c caaaatcctq | g actcaccata | aaccttgacc | catggtgaga | ı atgtcaatcc | 60 |
| ttaccctcgg | g aagcaaaaa | a agaatagagg | r ggaaatttco | aatcaaagaa | ı aaagagaagg | 120 |
| aaaatttcca | ı atgaaagcaa | a aaaaagaaaa | gaaggaaaat | tececaatea | aagagtggga | 180 |
| gaaagcaaaa | aaagaaaaga | aggaaaatto | cccaatcaaa | gagtgggaga | aagcaaaaag | 240 |
| aaaagaaagg | aaaattccca | ı atcaaagaat | gggagaaagt | aaaaaaggaa | gaagaagaag | 300 |
| gagagaaagc | tcctcatcaa | ggatcaaaag | aaaacagaag | atatgtgcag | agaggtcttt | 360 |
| ggaccggaca | atatctgaac | aatacagaa | | | | 389 |
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| | | atcaaacata | | | | 120 |
| • | | ttcaaacact | | | | 180 |
| catgtgtgat | gaaaaatttt | gccccaacat | taggaagtcc | aaattgcgtc | gcctcccacg | 240 |
| acacttccac | tggattcttg | tacacgcaaa | acaatgtctt | catcaattcg | cccaacggat | 300 |
| cagcttcagc | tgcaacctca | gcctcaccaa | gacgtctaag | tggcaaatgc | acattatgat | 360 |

| ttgaatcc | ta tatggtacaa gaacatcaga agttaa | 396 |
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| <223> <400> | unsure at all n locations 19662 | |
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| tgttccata | t gcgatccacg tgttggtact cagcctggac cccctcttat ttgtgcatct | 120 |
| ttatgcgag | a gaatttatga cgcatgctca aatgcttact tctctatgga tgtgaaaaca | 180 |
| caggtacat | g gatggctgtc acttttcaaa atcgtctgta caagattatg aaaataagtt | 240 |
| cagtgtcct | g catcagctgg taaagaaaca aaagtttgca tagttttggg agtcattttg | 300 |
| tttactctt | c tittccccaa cattcatttg ataagctgtt gtacagtgtt gacatgttac | 360 |
| aggattggt | t gtaataattc atacataat | 389 |
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| tgtccacaca | aacctgacct tatcattagt attaaagtat ataaatcaag gctcaggaga | 60 |
| gattgatcat | tttatggtgt ggctgcaaat aattaaccat tcaaaagttt ttaaaataaa | 120 |
| ctaattatgc | atgttcaatc aaaattttgt tgtctttcat aaaaaaaaat gttaatgggc | 180 |
| aaacataacg | gtatacacat ggggacaacc atttaagctt gatgtcaagc aaaacatcat | 240 |
| ggcttcaacc | ccacgatggg tcatgggtgc aacccgctgt ctcaagtctc tacaaacatt | 300 |
| tgcacagaag | ttgttcaatt cattgcattt ataattggga taattgtatc tttgttttta | 360 |
| atggatctat | ttcgaattga aagcttatta tatttggagt tcattaggac attacgcctg | 420 |
| tattgtat | | 428 |
| | 19664 390 DNA Glycine max | |

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| | att tcacctttat caatgttgga taatgagaat caaagcgaaa tcatcatt | |
| | cat attcatgatt atgggcacat gcatgccaag cttggagtca tttgtgact | |
| | ctg tocatgagat aatottttat atttotgttt tggtttatat attgottta | |
| | ac ccacaatttt taatccatta ttgcattgaa gtcaactatc ttattcaaa | |
| | aa ctaaactttg atctgcttct tttaagaata acagggaaca tgtatactg | |
| | tg tttaagaatt tattggggta | 390 |
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| tgacctatç | ga aactcagett eggattteta ggggeeetgt gtteegtgeg ttteeaceat | 60 |
| | g cgaagagcac ctttggtttt cccagcacct tgcactatat actcacccat | |
| | c cateetteae tgageteaeg tgeteeeaet gtteetatat eegtgataet | |
| | g agteeteata aateeteaca tgeett | 216 |
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| tgcttctttt | atgattagat tocattgcaa gcataaattt tttcttgaag aaacaactta | 60 |
| | aagtatttaa atagatgtat ccccatcctc aactatatac attatattca | 120 |
| | cttactatcc tgagaagtac tttttgaggg ataattaaag tacttacact | 180 |
| | cgtatctaac atcaattttc ttcaatgtga tttttccttt tttctttggg | 240 |
| tttatacctc | tttcactaat ttatgacatc tttttgtaca aca | 283 |
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| | | |
| ctttgcaa | ta aaatttottt acaaaattat aattaaaatt tagaactaca ctaaagatag | ı 60 |
| tacaaaagg | gt actttgatct ctaattattt aaaattcttt catctgatta tttaatccgg | 120 |
| ttgtaaaat | ca gatctggtgt tgtctttata tcatccatta aggaatagaa atttattcaa | 180 |
| tccataaco | gt tttggttgat aacaaatttc cttaatatac attagacaac agcggaaatt | 240 |
| gttttcaat | a tttgattttt cattatattt acaaaagtct aaacaagaag cattcttgtg | 300 |
| tgaatttaa | g aattttgaga tttcaaagta aaaaccttaa tatcagaagt taattatacc | 360 |
| gcanacctt | a attcatttat ggattangaa ggttcattag ttaatcaaat atccttacat | 420 |
| agatgata | | 428 |
| | | |
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| <213> | Glycine max | |
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| | | 60 |
| | cagatacaat actagaaggc attccatgca accttaccac ttccttgatg | 120 |
| | cgagtttctc cattctatac ttcatattca ccggaataaa atgagcagat | 180 |
| | gatcagetat gacceacaca geatcatgee caegaetagt ettgggeaaa | 240 |
| | aatccataga tatgctctcc catttccatt cccgaattta caatggcttc | 300 |
| | atggtcgctg gtgctcaacc ttagcctttt gacatgtcaa acatcttgct | 360 |
| acatatt | | 367 |
| <210> | 19669 | |
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| <213> | Glycine max | |
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| | | |

| atgcaaaaga | a tcaagaggag ttagtgggtt aaaaccataa a | acaacttcga | aaggagaaca | 120 |
|-------------------------|---|-------------|------------|-----|
| attagtggtg | r ctatgaacaa ctctattgta agcaaatgca a | acatggggta | aacaagcttc | 180 |
| ccaagtttt | aagttettee teaaaaetgt eetaageaaa g | gttcccaaag | tcctattaac | 240 |
| aacttccgtt | tgcccatcgg tttgtgggtg acaagtggtt g | gaaaataaca | atttagtgcc | 300 |
| caacttgctc | cacaaagtcc tccaaaaatg gcttatgaac t | tagagtccc | tatcactaac | 360 |
| aatgctcctt | ggcaaaccat ggagtctcac aatc | | ž | 394 |
| <210> <211> <212> <213> | 19670 358 DNA Glycine max 19670 | | | |
| agcttgaatt | ctaagagagc acaaatccta gacttaccca a | tttgtcttt | tcaatccact | 60 |
| | actagetttt caettgaett tgttttaaca ac | | | 120 |
| cttcttttt | tttaacttac aacattttat taattttttg to | gtgttctgt | tgtttcttac | 180 |
| ctttaaaatt | atccatcaaa ccaactcccc caaatttggg go | caaaattgt | cttctaacga | 240 |
| tgtgctctcc | taaaaccaaa gcatggtaaa tggagatgcc aa | attcaaagc | tcaaggttca | 300 |
| atttgacaat | tacaattcag ctcaaagatg ggtgcaaatg at | tatcatcat | tgagaaac | 358 |
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| | unsure at all n locations 19671 | | | |
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| aacaagtttt | ccacatccac aaatcacgta Laaacccacc at | cccttgtt o | gcccacctcc | 120 |
| aactgagctc | acgtactccc acgtagccca tatcctcgtt tc | tctcatca c | ccgagtcccc | 180 |
| ataaatcctc | acaagettee ecaacateea ggtaatteaa ca | itccaaatc a | itcacaaact | 240 |
| aaaaaaccaa | gcaaaacagg gcaaaggtag aaaactctgc cca | aaaactca a | accaaaatc | 300 |
| acagcttttt (| ctcacttaaa gaccccagta acatttcctt cg | ttccaatt c | gttcaccgt | 360 |

| tggatcgac | t cgaaaatttt | actggaagto | tctagtacat | aagcctacan | tttgacc | 417 |
|------------|--------------|------------|------------|------------|------------|-----|
| | 3 | | | | | |
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| <211> | 399 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | ζ | | | | |
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| agcttgtttg | g acagccaatt | tgaggggaac | tgagcggttt | tgacgcttca | ttaaaaattg | 60 |
| tcgaactgaa | ccacctttgg | catactcaga | tacaatgcac | cataccattg | gcttgcggca | 120 |
| tgcaccaatg | aaacgaacta | tgttagaatg | ctttagtgtg | gccaacattg | tgacctcctg | 180 |
| ctggaactgt | tgttccatca | attgagcctt | tgctggatca | ttttcaggcc | tctccaagat | 240 |
| tttgattgca | acatcttcac | cattgtaagt | acctcggtaa | agtttcccaa | aagctccttg | 300 |
| agcaaaaggc | tcacccatat | tcagtttcct | gatatcaatt | gtccactcat | caaaattgtc | 360 |
| aagcccttca | gtcggagaac | tattgtccat | tatagcttg | | | 399 |
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| <211> | 449 | | | | | |
| <212> | DNA | | | | • | |
| <213> | Glycine max | | | | | |
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| gattgacaca | ggcgatgaac t | ccttgtcac | tgaattaatg | tttaatggtg | aatcaatatt | 120 |
| tgtttcccaa | ttatagttta d | catttcatta | tgttttgtaa | cctgtttttt | tcccttcatt | 180 |
| tcccctccac | cattaggtac t | tttaatgac | cttgaccatc | atcaagttgc | tgcccttgcg | 240 |
| agttgtttca | taccaggaga t | aagtcaact | gagcatatac | aactgagaac | agagcttgca | 300 |
| aggcctctgc | tacagettea a | ıgatagtgca | agaaggatag | ctgaggtagg | tgtttgttca | 360 |
| cttaacctga | atgtatcttc t | gaattaaac | tataaatggt | atcacccttt | tgtcacagat | 420 |
| acaacatgaa | tgcaaattgg a | tataaatg | | | - | 449 |
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| <211> | 401 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |

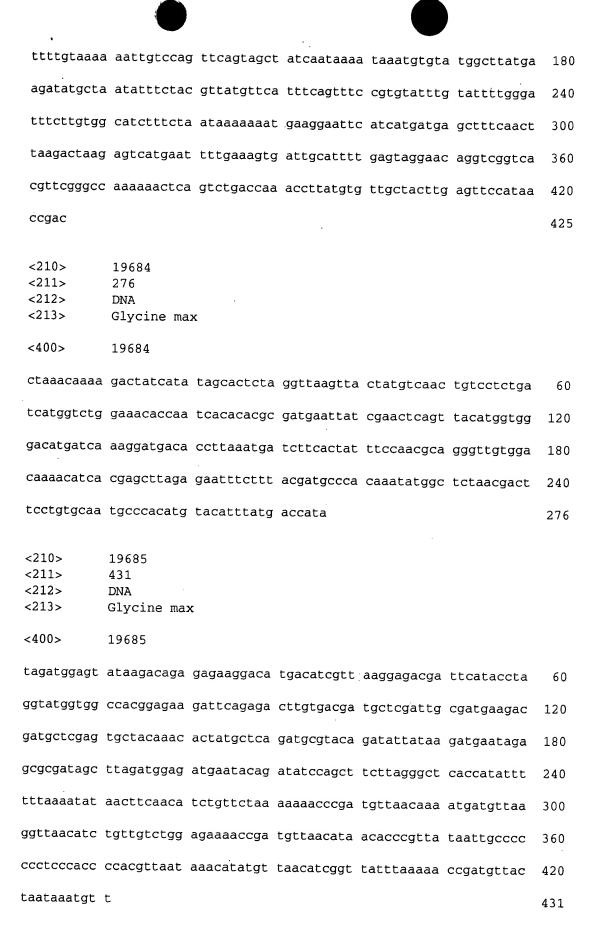
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| cttgaagcaa | ı tgttttgaat | gtttattgaa | gtaatcttga | aagcaacctt | gtttgattat | 120 |
| tctttgacat | catcaaaato | atgtattcat | acattcacat | atactaattt | gagtcttggc | 180 |
| cacatcgtct | acaatagtta | ı aaggtaactt | acatttaata | aaaaatgtta | atgagttaga | 240 |
| taaaaaaact | taaaagaaaa | ı attttgaaaa | ttttaggaac | caaatgtaat | aaaattattt | 300 |
| ttggaagact | aaaactaata | ı tgagtcaatt | aatatttgga | gtaacaaaat | catatttaac | 360 |
| tcttatttat | aggtctatca | gtttattatg | tgaccaatgg | a | | 401 |
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| tatcatcatt | ttttctccgt | cattgaggtg | ccacttgaac | tgctaggtct | ctccaccttt | 120 |
| gggcgtattc | ttttgaaaga | ttcgtgcccc | ctttttgcac | atgttctgta | gttgcatcct | 180 |
| atcctaagac | atcatattga | cactgcttaa | cgaaggcaac | cactangncc | ttccaagaat | 240 |
| ggactcggga | aggttccaag | ttagtgtacc | aggtaacagc | taccccagta | agactttctt | 300 |
| ggaaggaatg | tatcagcaat | tcctcatctt | ttgcgtatgc | ccccatcttt | cgacaataca | 360 |
| tctttagatg | gttcttgtgg | caagtagtcc | ccttgtactc | gtcaaagtct | agcaccttga | 420 |
| acttgggagg | ggtgatgata | ttgggtact | | | . ' | 449 |
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| gtggatggcg | cctcctctca | cctcttctct | tttgtcttcc | gccgcattga | aagctcattg | 120 |
| aagctcattg | aaggacctca | ttgaagctca | aagatccagc | ctccatagaa | gctccacaag | 180 |

| caagettee | a tcaagtggta | atcagagcat | aagagcțtca | a agtaggtgct | ccttaaacca | 240 |
|-------------------------|------------------------------------|-------------|-------------|--------------|------------|-----|
| tgaattgtgt | tgggtttagg | ttcctttgtg | , tttagtttt | c atatagaago | tagatttgat | 300 |
| tctctatggt | tcatatttct | tgttcttgtt | cttgaaccat | gaattgtgto | tggtttaagt | 360 |
| tcctttg | | | | | | 367 |
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| tagccgaatt | cagatcgaat | tgaagttagc | ttagctcatc | cttggtcagc | ttagcggacc | 60 |
| aaatcagcct | tagatgcaag | ggttgggcac | taagcgcttg | agactcgcag | cttagcgcat | 120 |
| gaacaaagat | gcgcttagcg | tgaggcttgc | gcttagcgaa | aggactactt | tttagaaaaa | 180 |
| agttttctaa | gttatttttc a | agtccttttt | ccaaaaaatt | gaaaccctta | tgttaaacat | 240 |
| tcaaacatag | gctaatatac (| cctatgtat | agatcataca | acaagttcca | aatgattaaa | 300 |
| tgcattaaaa | acaaagataa (| cagaaattaa | aaactgggtt | gcctcccagg | aagcgcttct | 360 |
| ttaacgtcat | tagcttgacg (| cttttacctc | accgggtgat | cttatgtttt | ggttcttact | 420 |
| ttcagaacct | cttgacct | | | | | 438 |
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| cctaccttgc | aagtattttg t | tcaataatg | tgattgaagc | aaattttta | ttagatttaa | 120 |
| ttaaataaat | gtttagtatt t | tgttctttc | ctattagtct | gttgatagct | aactggaata | 180 |
| aaagaaaagc | taaggtcgga c | cagttgata | gctaattgga | ataaaagaaa | agctaaattg | 240 |
| cagggaataa | ttatgatatc t | tttatttc | atcattaccc | ctttntatag | ccatttcata | 300 |
| caagatattt | tgctaagttg t | tataacaga (| attttgaaat | tgcataacca | cacaggtatc | 360 |
| | | | | | | |

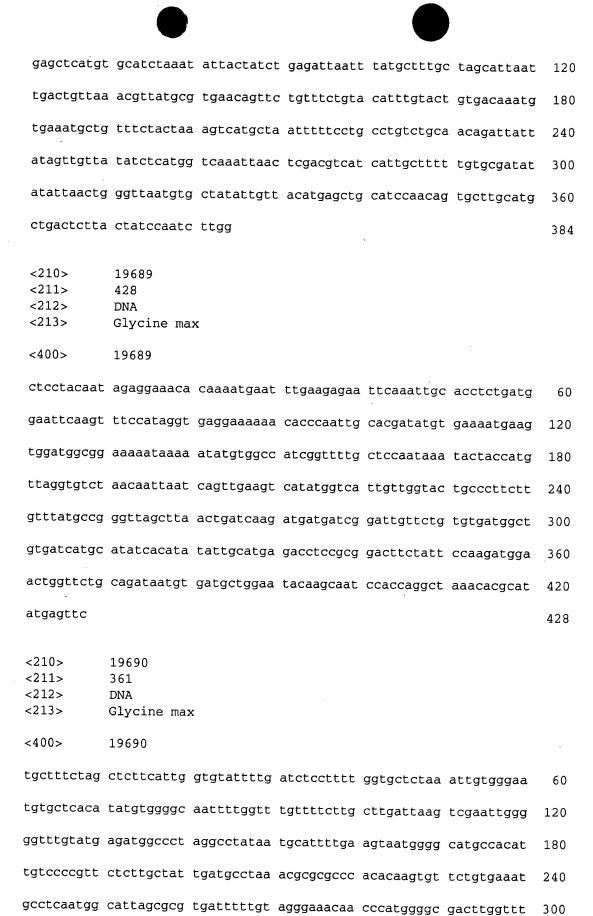
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| gttttaagca tacgaagcct gttggtcact atctcattga ta | aaaacttgg aggttaagag 120 |
| ttgaaggtct agtagtgaac aatcacgctg taataatcaa ta | aatggtccc cacccctctc 180 |
| ctattattat tattccaaaa gagctacttg aatcatcact go | ctcaaaaat tcagagctgg 240 |
| ataagttgta acaactgaca aactcttgta ttctctccat ct | ttgaaccct acacaccact 300 |
| gtgtgtagtt gcaacttgca agcttgattt gacttcatct at | cactactat atatgatttg 360 |
| cacaatcact ttagctcgta tactagttgg ataggtgttt aa | attctagcg attagtcaac 420 |
| aagcacg | 427 |
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| tctattacta ttcctgttta aaaaaatttc atgcggaaag tt | gaattcta aggaaagaac 120 |
| tctcaagaat tttactttta tcaaaaaagt attattttt aa | tgtcgttt gagggaaaag 180 |
| aaagatetta tttttetttt atgtacaagt aaaatecaaa aa | tccaaatt gaattttatc 240 |
| agtaaagtaa acatgataaa agttctttat catgagacct ata | aaaaaata atctaaaaac 300 |
| aattetteea ttaaaatget ettaatgtte acatggegag aca | agagaagt tattaacaaa 360 |
| cctttgatat taacaatgaa gttaccaatc a | 391 |
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| aaaacagago | c aaaggcagaa aactctgcca aaacaccaac caaatcacag cttttctcac | 120 |
| ttaaagacco | c cagtaacaat teettegate caattegtta acegttggat egactecaaa | 180 |
| attttactgg | g aagtetatag tacataagee tacattttga eegttgggat etactageaa | 240 |
| acattcagaa | a ctcattctgt actactcttt ccacaaccaa tcacacacaa gcatttttct | 300 |
| gcacaaagco | c aaaatcctgc tgcacctatt ttgacagcaa aattctgcat aagtgcagat | 360 |
| ttcgaaaatc | c açacttgccc tcatccaatc | 390 |
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| | 19687 | |
| | agaagatett caaagagatt teaaggettg taaaggattg attggaaaag | 60 |
| | caagattgct gttagaaaga ttgattggaa aatgaaaaac aaagccttgc | 120 |
| | tcttcatgtc tggtcaagaa ggtcattcag aagagttata acttttagaa | 180 |
| | cccatttgaa aaagtcaaaa cctttttgaa gagttacatc tatagatttt | 240 |
| | acactggtaa tcgattacca aatatgtgta atcgattaca caaagctttt | 300 |
| | atgtgactct tcacttttaa atttgaattt caacgttcaa ggacactggt | 360 |
| aattgattac | caaaacattg taatcgatta cagccttttg aaaatatctg gaacatt | 417 |
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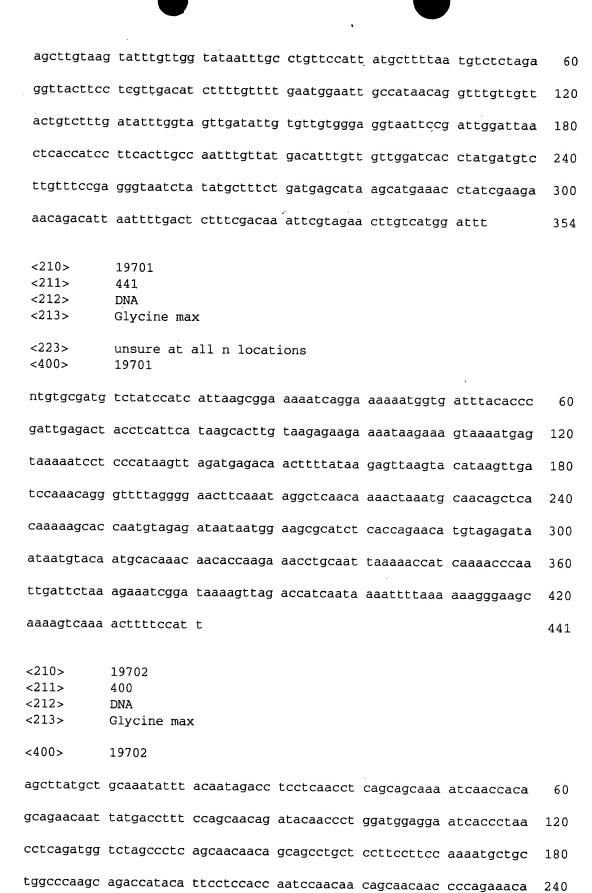


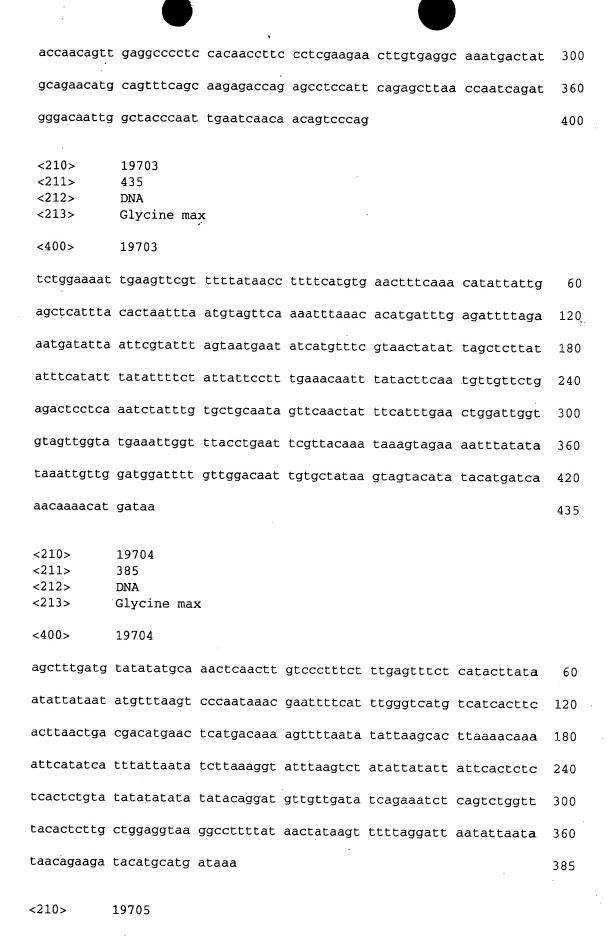
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| | cacatgette | tatttataga | ctatgtagct | tccttgataa | gctttcttga | gaaaacttcc | 180 |
| | ttgagaagct | tctttgagaa | aacttccttg | agaagctaga | gcttagccac | acacacccct | 240 |
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| atctccaago | : cttccttcca | a ctcttcacca | a tttttgtgca | ı ccaactgag | y tatggaggag | 240 |
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| gcacgcgctg | tctgtggtct | tecetetegg | catcgagccc | acactgttga | ctgtgggtgc | 240 |
| tgaaactgcc | gccactacct | gcaaggtatg | tgaggtttgc | agcgacggca | ttgtgactga | 300 |
| ccgcggcgag | gccatgtcac | tctcctccat | aacacttcct | ccgatgaact | a | 351 |
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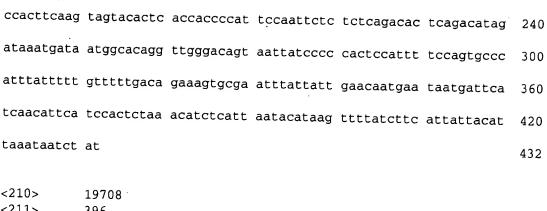
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| | | i.X | | | | |
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| tttttttata | tattttccat | tgaatgctct | tctcccccat | : taagcttgtt | caaatttcta | 180 |
| tccattaaaa | gcagcctcct | ttaagtagac | ttttcatctt | agtaatagga | aatgaaaagt | 240 |
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| ccaaaaccaa | gcttgaccaa | tcccgaccca | acccgggcat | agtcggtcag | tgagaacctg | 180 |
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| aagcaaggag | gcttgtggtg | gctggccagc | tgtgaatttt | gtgtaatatg | tggattgtgg | 300 |
| cctctggtaa : | tcgattacta | agggtgggta | atcgattaca | aggcttaaaa | ttgaggacag | 360 |
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19708

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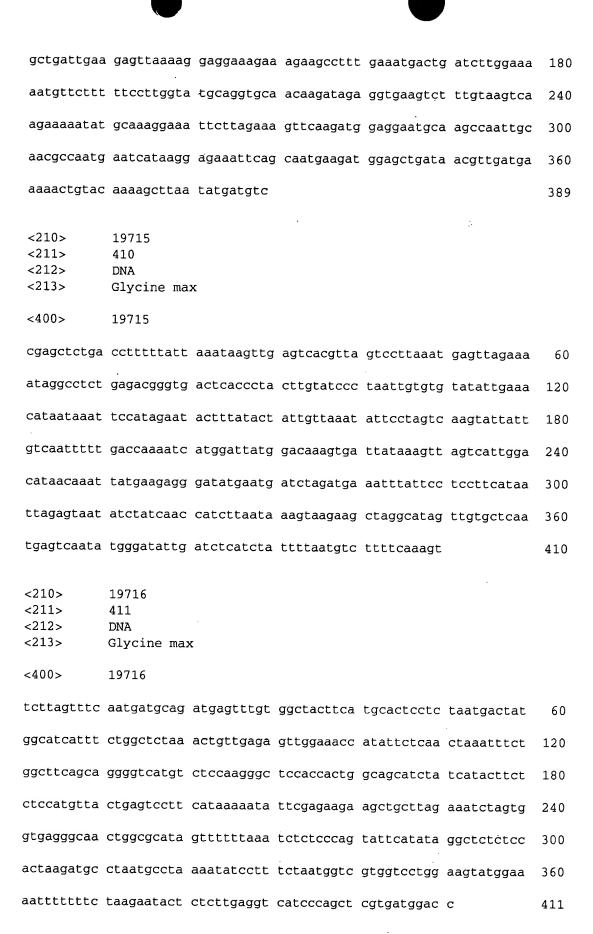
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| aagaaagagc | agaaggaaaa t | tccccaatc | aaagagtggg | agaaagcaaa | aagaaaagaa | 180 |
| agaaaattct | caatcaaaga a | tgggagaaa | gtaaaaaagg | aagaagaaga | aggaaagaaa | 240 |
| gctcctgatc | agggattgaa g | gaaaacaga | agaaatgtgc | agagaggtct | ttggaccgga | 300 |
| caatatctga | acaatacaga a | ttgtcacca | aatgaacaaa | aaagaaggaa | agganaccac | 360 |
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| actttcaatc | tgcccctaga t | tcgcttgag | gtttatgcat | ggtacctttc | attgccccag | 120 |
| tgtagggctt | tgaggtatcc a | tcgttgttt | tgttttcaca | accttgtagc | aaggaagaat | 180 |
| gaaagaggcg | gtttgattct c | gcaaaaaga | atttttcaag | gacgagaaat | agttgaagga | 240 |
| tttttttga | gttgacgggt ta | aagtcaaat | gactcctatt | cttgataact | cacttctctc | 300 |
| taaaaaagac | aaacttttag ga | aatgataaa | atgaggtcac | atgaatgtct | atatttttac | 360 |
| ttgaaaacac | agtcaatcaa at | tgctntttt | ctttntcttt | gtgaactctt | tttttttgc | 420 |
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| taatataata | aataatactg acgtgaccaa acaagaaata taaaaaagtg tcaaataagt | 120 |
| gcatctaata | acagtattta gacttttcct ccctgattta cttaaatcca cgtgattttc | 180 |
| attattatct | taatatcagt atctaattac actgtttcct aaaggcatat aacattgtgc | 240 |
| accaacttct | gcctcatcga ttcccaacta tccaggggac gttctgggca acaattttt | 300 |
| ttatttggtc | aaattaaatt tcaaatgtcc acacctagca tgtagattca gggtcaagaa | 360 |
| ggaagttgtt | tatatactgg taattggg | 388 |
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| gagtctgcta | tetteaateg aetttttgtg gagegggeat acaagtateg eeetgttaag | 120 |
| gtggtggaat | ttgaacttcc gcggcagcag tgtgtggttt acttggatct gaagcgggag | 180 |
| gagtgcacca a | atttgttccc atctggccga gtatattcac aggcattcca tttaggtgga | 240 |
| caagggtttt (| tettateage acattgeaae atggaceaae agagetettt ceattgettt | 300 |
| ggcctgtttt t | taggaatgca ggaaaagggc tcagttagct ttgccgttga ctatgagttt | 360 |
| gctgctaggt (| caaggccaac agaggaattt gttagcaagt acacatgcaa ttatgtattc | 420 |
| ac | | 422 |
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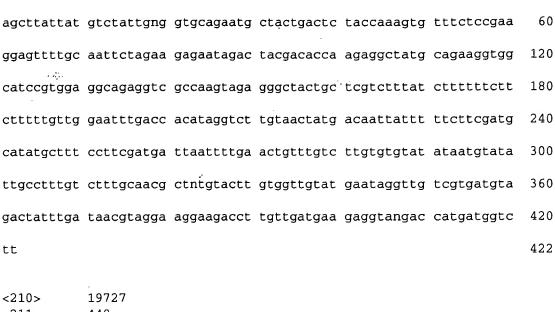
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| tctctttgga | a aagaaatete aaaactetae aaacaaagge accaataage tttettetgt | 180 |
| tggtgttgtg | g gaggaattat ttgaagaggg ttctgcaatg cttacagaga ggtaactggg | 240 |
| accataacto | g ctaaatttat atttgcatga tgtcatttga agtttaattg gtcaccatca | 300 |
| tggtggaaag | g agagaaaaaa tgantccttt ttcccaatac atctatcttt gattctntaa | 360 |
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| ataagtgtta | ccttgtatgt gatgttaaga acataattag ttttactaat ttgtaggtga | 120 |
| atctcgagtt | gtacgtactc ttgctttcac gatctatctt ctttttgcca tatgatttta | 180 |
| gacaaaagta | taagttggtt ataaaggagt tttccgcatg gtctcgtggc ttttctctat | 240 |
| agaaaaatac | agcatagaac ctgcgttctc gttaaactgt actattttca ctactaagga | 300 |
| catgattacc | attgtgtttg gttgaaaggg agaaaaagaa gtgaaatagt gaaagagagt | 360 |
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| actcancco | g aatgacaaac cggaccaaag accccagcac cacagattaa cccaccccca | 120 |
| naggggggg | a gaacaaacga cccaaaacac caaaaggaaa agacaaacag aaaaacgcaa | 180 |
| aaaaaaaca | c aaaccaagaa caaacaaaac aacacacacg acaaaaagaa acaacacaac | 240 |
| aaaacaaaa | c aaccgaaaga cgaaaaaaga ccaaaaaaag aaccacaaaa caaaaacaaa | 300 |
| ccaccaaca | a acccaaaaca gacaaaaaca caaacaaacc aaaaacacac accccaacaa | 360 |
| caaaacaaa | c cagaacg . | 377 |
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| | c cacgatggag atgcagcgga agacaaagga gaagaggtga gaggaggcgc | 120 |
| | ggaataagcc atggaagaag gagcttcacc accaagatga gccttggata | 180 |
| | agaggatgct tcaatggagg aaaagaaaga gggagagaaa gagagagggg | 240 |
| ggagcacgaa | attgaaggaa gaaaaaggga gagaagttga actttgagtt gtgtctcaca | 300 |
| agactctcat | tcatcanagt tacaacaagt gttacacatg cttctattta tagactangt | 360 |
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| agctcaccac | cataggagge catgaataag agettggagg aagaagaaga tgaatgaagg | 100 |

gagaggaaga gaagagcacg aaattttgtg ctctaaaaga gctataaaat ctgaagttta 240

| attttcaaat | gatcaaagtt | gaaaaaatgc | acacacatgg | tctctattta | tagcctaagt | 300 |
|-----------------------------------|------------------------------------|------------|------------|------------|------------|-------|
| gtcacacaaa | attggatgga | aatttgaatt | tctattcata | tttcacttga | atttgaaatt | 360 |
| aaatntgtgg | agccaaaatt | tcactaatta | tgattagtga | attttagcta | tggttcagcc | 420 |
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| tcttaaattt | ccatggcata | cctaccaaca | agcttagcac | tcattaacct | tggtggagtc | 120 |
| agccaagatc | ttttattaaa | aatggtgaca | aaaaatcctc | cttgacttga | gtgaagccac | 180 |
| cgatcactac | atgtaatgac | aatgttgtcg | acactcagca | tagataatga | agtaaatgca | 240 |
| cccaatagcg | aggtggcgat | atataatgag | gtaactatcc | tcactctgag | tcataccaaa | . 300 |
| ctattgaaca | atggtgctaa | gtctacaaat | acatgctcta | gagattgttt | aagaccactt | 360 |
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| ttatta | | | | | | 426 |
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| cgttatgtat | gccactcaaa | acgacatgta | tttcaaatgc | caagcgaact | ctgatacttc | 120 |
| gatccatgaa | agtcattttc | taagccgccg | tcaagctatc | agttccatat | agattacagc | 180 |
| tccatcaagc | tgcgtttcaa | taataattct | acacatactt | ttgcaaataa | tatgtcttaa | 240 |
| cttacaccaa | ccacaaccac | tacatattca | attacgaacc | gcatatatat | aatctcgtta | 300 |
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| gatcattgac | aatgtttcat aatgcagaat aatttattct tttgcagcat tgtgattttt | 120 |
| caatcacact | tggatttgga taggttccaa ttaaggcaaa aattattata tttgcttgat | 180 |
| caactaaaat | gttctttgta catatttttc tgtgtatata atattaattt atgtatatct | 240 |
| aattttaat | atttctgtta tttattgtta ttgtattttt ttaattatca tgtgatgtct | 300 |
| tgggttttat | ttgttaggtt tttttatcat tctaatcaca ttggtgatga tgttaattta | 360 |
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<213> Glycine max

<223> unsure at all n locations

<400> 19727

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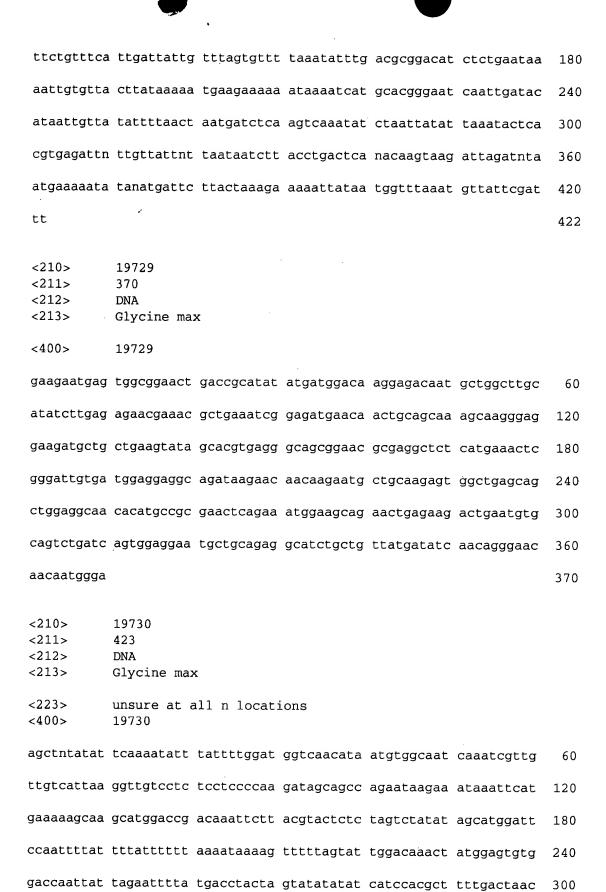
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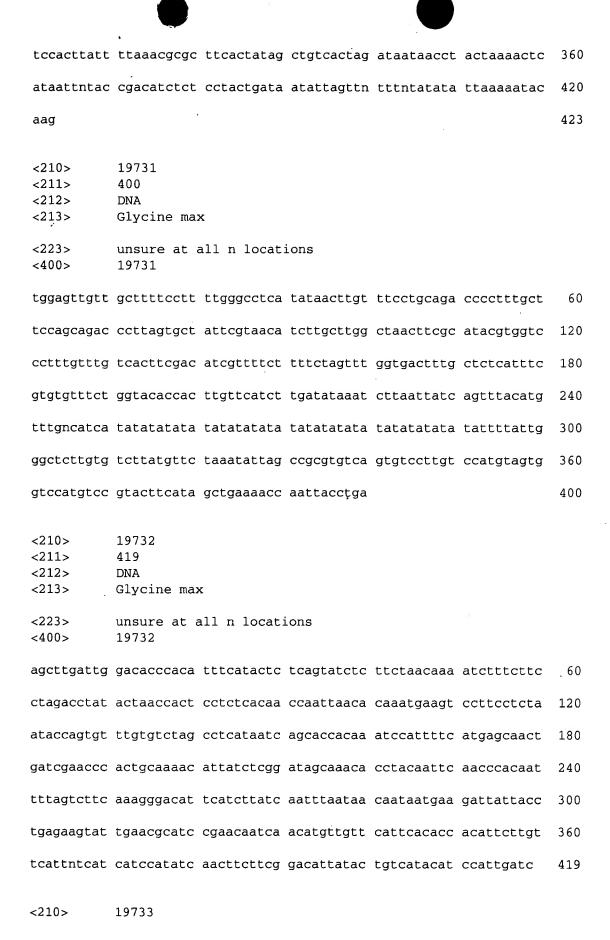
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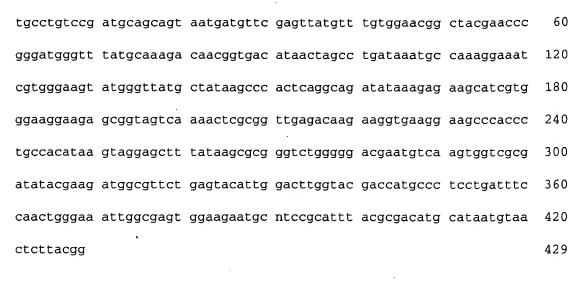
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| attaaagagc | tctaaccaag aacatgggaa aatcaatcat tataatcacc acacatccac | 180 |
| cattacaagc | atgtaagtet teetttgege egaacacaat teatatgaag aactgeacae | 240 |
| aacatctatg | ttgattcaat caaattaatt aacctacaca atagatgtta ctttggtaat | 300 |
| atgctatgtc | aattaatcca accaaaaata ctagacaatt tagttaaatt taaatttaca | 360 |
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| | | |
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| <212> | DNA | |
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| aatggtgcta | attatacaaa tcatagagta atccaaatta atcatgctta aagtaaatat | 180 |
| acacaatgta | aatctacatt gtctaatctt gcaacgtgta attttacacc gtgtagaaat | 240 |
| | | |
| caggttctaa | tgaaattcta atttctacta aatttatcac atgaaatagc atgcatataa | 300 |
| | tgaaattcta atttctacta aatttatcac atgaaatagc atgcatataa aattggttac ttgaattaaa tcttccttat aattactttt aataccaatt | 300 360 |
| tcggaatttt | | |
| tcggaatttt atatntccac | aattggttac ttgaattaaa tcttccttat aattactttt aataccaatt agatcaccca aactcaatac tttctgtaat | 360 |
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|-------------|-------|
| <211> | 419 |
| -212 | עזעכו |

<213> Glycine max

<223> unsure at all n locations

<400> 19736

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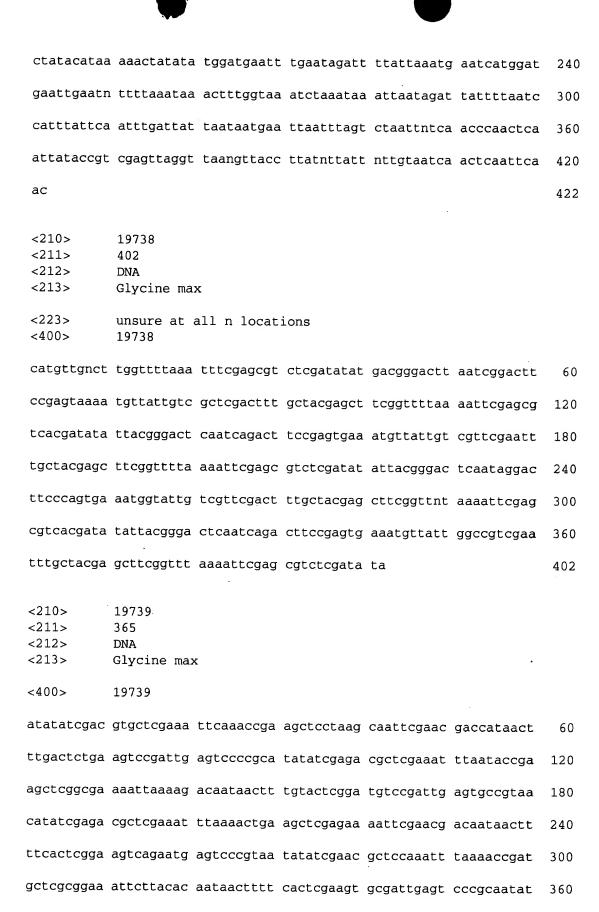
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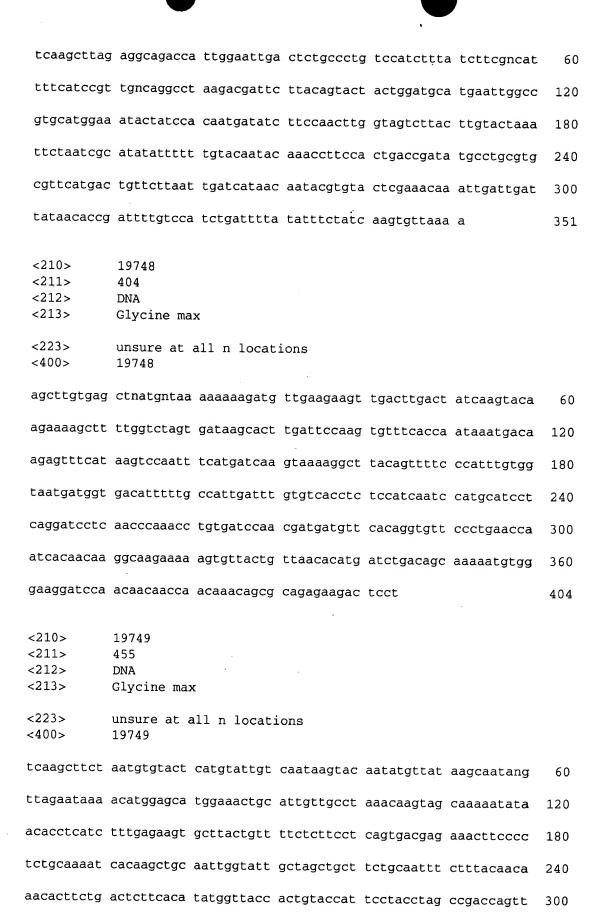
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| tatattagct | gotttttttt tttttttgcc tctcgtccaa gtatttatat tcattattta | 180 |
| aatcgacact | tatgatcgag gcatatttcg ttaactttta tacaaggtat gactgatatg | 240 |
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| tcagtttgaa | ggagaagtaa gctttttca tggcactgct ctttatgttt tttctcttgg | 180 |
| ctttgagaaa | atactaagta tttgaattct tcagaggcaa gatccttgaa gtactgaaaa | 240 |
| actggccaga | aaagagtatt caagttattg ttgcgactga tggtgagcgt atattaggac | 300 |
| ttggagatct | tggttgccaa 🎉 aaaatata gtgttagtct cctatatctg cattacacac | 360 |
| agagacatta | gagtaacata attttt | 386 |
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| agcgtacgat | atattacggg accatcagac ttcagtgaaa gcattgtcgt caaatgctac | 120 |

| agcttcgttt | aaatcagcgg | g cgaatatacc | ggaccataga | ttccaggaag | gattgccgtt | 180 |
|----------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| cacttgctca | cctcggttaa | attagcgacg | aatttacgac | tcataactcc | aaggatgtat | 240 |
| gccccaatac | acgacctggc | : taaatgacgc | caatatac | | | 278 |
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| cgcttcttta | acgtcttgag | ctggacgcgt | gatgacttgt | cggccacgga | cctagtactt | 180 |
| tgcttacctt | tggctttgga | cttggtcgcc | tgctggtcga | ccacgggtcg | taggcaacgc | 240 |
| tccagccttt | gtagatgagc | tgagggactc | tggaggtggc | ggcgatgcgt | ctattgcccg | 300 |
| ctgccggcca | tccccaagct | actgtggtgt | ctcgccttgc | gcctgcctgn | gggcgcagta | 360 |
| cttcttgatg | aaagctcggt | tagtaggggg | cctgatgacc | ttgatgtggg | cgacgggcac | 420 |
| tccgtagaac | tgacagaggc | ccgtaatca | | | | 449 |
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| agctttcaat | gaactcttgc | tccaaaaaat | tgttattcga | tccaattcaa | aaatcttaca | 60 |
| aaatggtcca | tcaataagca | tgtgaaatct | aacaaaataa | caagttaagg | agaacaaact | 120 |
| acaacttaat | aacttacatc | ataaaaggca | taaacaaagt | cctaaataag | agaaaagatt | 180 |
| agataatttt | ctaaattcac | atgtctcagt | taagtatttt | tggcaattat | taactccccc | 240 |
| aactttagaa | atttttttgt | cctcaagaaa | aagtaaacac | attgttaaat | gaaaactact | 300 |
| agtgctaaga | ctaaaatatt | gacatgagtt | gagatcaatt | ccatcctaac | atatccaaca | 360 |
| cttgtattgt | ccagaaccaa | ggtcataaaa | aaggaatggg | acaaggactt | aattattctc | 420 |
| agtgatgtca | ttg | | | | | 433 |

| <210> <211> <212> <213> | 19745 417 DNA Glycine ma: | × | | | | |
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| tttaaaatca | actttacgcg | taactttatc | gcttttttaa | gaactatata | ggtctgagtt | 120 |
| cctcttcgca | cttgaggata | cgtaggagca | agggtcatgc | tcttgtcgac | cccaaaagat | 180 |
| aaaaaacaca | aaaaagggaa | aaataaataa | atattgaagt | catgattttg | cacacttgat | 240 |
| taaaggtcgc | cgtcccttgt | gacggacgaa | tagggtgcta | atacctttcc | ggcatgtaaa | 300 |
| caactcttga | acctttattc | ttaaaattcg | cagacccctt | tntagttttt | ctaacgttnt | 360 |
| cctcgaataa | acattggtgg | cgactcccgc | atgtcttcct | tctttggatg | acgcacc | 417 |
| <210> <211> <212> <213> | 19746 412 DNA Glycine max | × | | | | |
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| agcttagcta | ttggtctgag | agatcccatg | accttctaaa | ccccacaata | gaatcccata | 60 |
| ctttcaagga | taaatatcac | ttctgttttt | tcgttttggt | gcattatgtt | ttaaaagtct | 120 |
| cattgcggct | atatgtttaa | ttgatgttaa | agaacacatt | ttacattgat | atccatttga | 180 |
| aaggtaataa | aatgcattgc | ctatctccgg | gagaaccaac | ccagtcagca | tctgagtatc | 240 |
| caactacctg | agcatgtccc | ctatcttcat | aaataaggac | ttagccagga | gccttcttaa | 300 |
| | | | | | | |
| tgtatttcac | ccagtgatct | tggcagggag | agtttatgaa | ctgacttacc | acactcactg | 360 |
| _ | ccagtgatct atttgggcaa | | | _ | _ | 360 412 |
| _ | atttgggcaa 19747 351 DNA Glycine max | ctgatcatga | cataattcaa | _ | _ | |



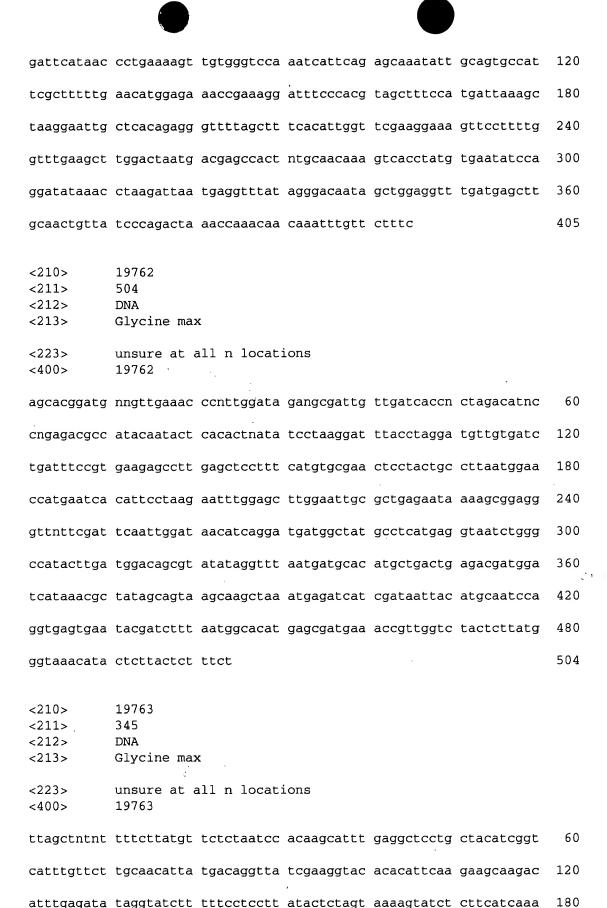
| cggccagctc | ggtgtgcata | gtgaattgaa | tctgtcggta | agtctagatt | aaccacaaga | 360 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| tcacattctg | ccacatccaa | accccttgct | gataattcat | tcgtaaccag | aactctcacc | 420 |
| tcaccattct | tgaatgtctt | cagagttgtt | gacct | | | 455 |
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| agcttcactg | tttattatag | tatgtacaaa | agagttgcag | tattcaactg | cactcttagt | 60 |
| taaagggagt | tagttaaaag | atagtttgga | agcttgttag | tcttcctagt | tagttcaaag | 120 |
| ttagctacaa | acagtttcag | taactgtttt | atatatatat | atatatatct | ttgtaacaca | 180 |
| atattcactg | gccaaatata | tttttccttt | cttgattctt | gagttttcct | ctctctcaaa | 240 |
| ctctctctaa | tcttctttta | tgatactgtt | tagtttctaa | taatgtttat | taatgataaa | 300 |
| aaagtgctcc | atgcacccca | tcagatttgt | cctttggtct | tcttccanaa | tgcttcaatt | 360 |
| ccagcccana | atagctattg | ggccaatggg | cccaagtatg | gacacttcag | cacctttntc | 420 |
| tttacat | | | | | | 427 |
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| <223> <400> | unsure at a | all n locati | ions | | | |
| ttacgcgacc | ttagaatcct | aagcnttcta | atgtcttgtt | agagtcgtgg | ccttcttcag | 60 |
| tttgtcatct | ttagtgtctt | catagnttat | gctcaataag | ttccaaacct | ctttagcagt | 120 |
| tctcagtctg | taaatcttgc | tgtaatcatt | ctttgatagg | gcacatgtta | gagtatatct | 180 |
| tgcttttgtt | ttaagttcca | taataaccag | atcatcatct | gtccattcat | cttcaggttt | 240 |
| atgaataggg | atatccctat | tggtaatcat | gagctagatt | tcatactggg | tggacttgat | 300 |
| gtacatctcc | attatgtcct | tctagtaagg | gtagttntct | tcagcgaagc | ctggtggcct | 360 |
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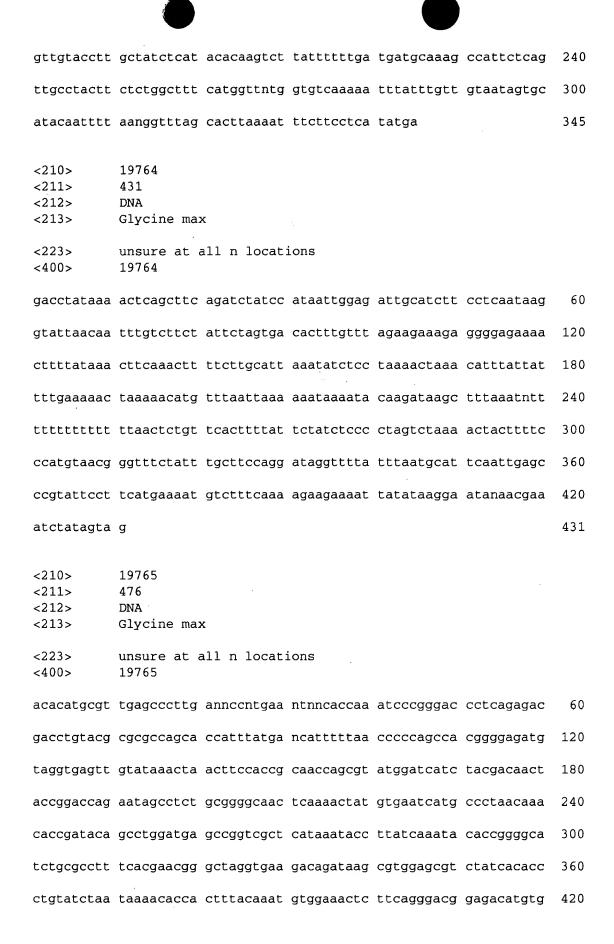
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| <210> <211> <212> <213> | 19752 . 432 DNA Glycine ma | ı. ax | | | | |
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| tactagacaç | g tcatcttcac | cttatatcag | g aaccagcaac | tgctaggtag | gacccactgt | 120 |
| ggtcgaacto | cactgcaaga | cacacagaat | gaagatgaag | gccttttgaa | atatgtacaa | 180 |
| cttcttaacc | agtatatata | aatataggtt | atattgccat | caagtgaaaa | tgaataccgg | 240 |
| agctcgttgg | , tgtttccgaa | tcatatggag | caaaattcct | gaagttette | agtttacgta | 300 |
| gatcccatag | cttaacacca | tcatgagcag | cagtctgtca | aatctaagag | tacaataaga | 360 |
| atactgtata | aacaatcatc | atactttact | agttacattg | tctcacatac | cgcaaggaag | 420 |
| tatccattct | ca | | | | | 432 |
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| agcttgcatc | atatgctatc | gacaataaca | tttcactcgg | aagtccgatt | gagtcccgta | 60 |
| atatatcgag | acactcgaaa | tttaaaaccg | aagctcgctg | cagacgctaa | cgacaataac | 120 |
| atttcactcg | gaagtccgat | tgagtcccgt | aatatatcga | gacgctcgaa | atttaaaacc | 180 |
| gaagctcgta | gcaaattcta | acgacaataa | catttcactc | ggaagtccga | ttgagtcccg | 240 |
| taatatatcg | agacgctcag | aatttaaaac | cgaagctcgc | agcaaatgct | aacgacaata | 300 |
| acatttcact | cggaagttcg | atggagtccc | gtaatatatc | gagacgctcg | aaattaaaac | 360 |
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| <210> <211> <212> | 19754 415 DNA | | | | | |

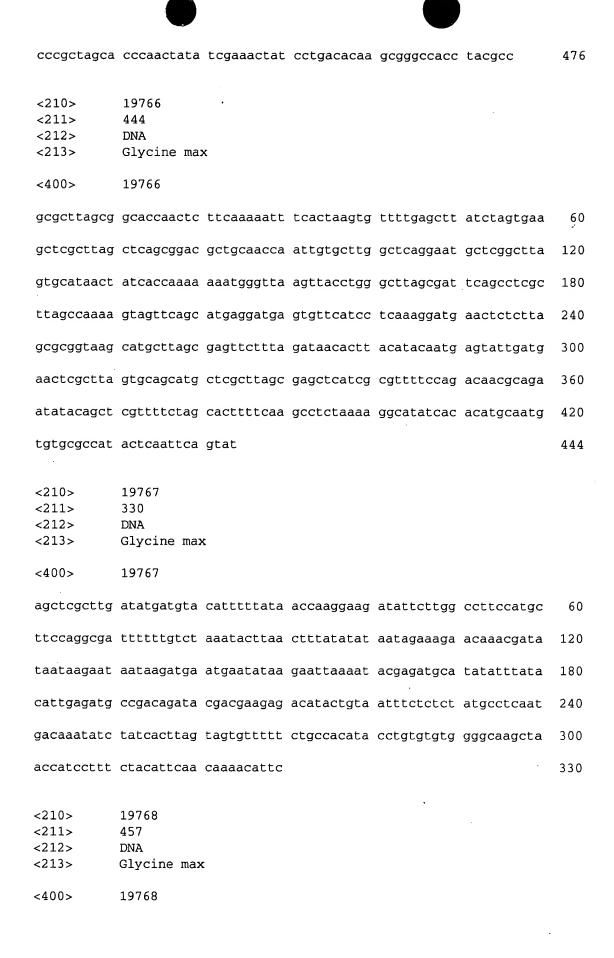
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|-------------------------|--|-----|
| <223> <400> | unsure at all n locations 19754 | |
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| gaatttgcta | a cgagettegg ttttaaattt egagegtete gatttattag gaeteaateg | 120 |
| gacttccgag | g tgaaaagtta ttggcgntcg aatttgctac gatcttcggt ttgaaatttc | 180 |
| gagcgtctca | a ttatgttacg ggacttaatc ggacttccga gtgaaaagtt attgtcgttt | 240 |
| gaaattgcta | a cgatcttcga ttttaaattt cgagggtctc aatatgttac gggactcaat | 300 |
| cggacttccg | g agtgacaagt tattgtcgtt cgaatttgct accagcttct attntaaatt | 360 |
| tcgagcgtct | cgatatatta cgcgactcac tcggaatttc gagtgaaaag ttatt | 415 |
| <210> <211> <212> <213> | 19755 376 DNA Glycine max | |
| <400> | 19755 | |
| agcttgtagg | ttagcttgca ggaatccttc gaattggcca atttcacttg gtgctacaaa | 60 |
| gcagtgacac | tagtcgcaga ggtaagggag gactaatatt tcaaaacatc atccctcacc | 120 |
| cactcatagc | aagcaatagc aggggggacg atagtgaccc aacacactat ggctacgcca | 180 |
| cactcggggg | gaageettaa egteeetgee accaettete ettgetttte tagaaataga | 240 |
| agggggaaga | atcgtggtta ctggaacggg ttccctgttt agagaagaag tcaaaggctg | 300 |
| cgccaacacg | gtgtgagacc gtcaacctct ccaaggaaga ggactcgcta tcagaaaccc | 360 |
| tttggcgagt | ctcctc | 376 |
| <210> <211> <212> <213> | 19756 242 DNA Glycine max | |
| <400> | 19756 | |
| | cataagccca actccttatc aaggaaataa tccaaccaga atttcaataa | 60 |
| cctaaaatgt | tcacaaccac aaaatattcc agactggaac acaagaaaaa taagccaagt | 120 |
| tcttatcata | attatggaaa ttctaagaaa ctaaaaagcc aaatacacgg cttataaaag | 180 |

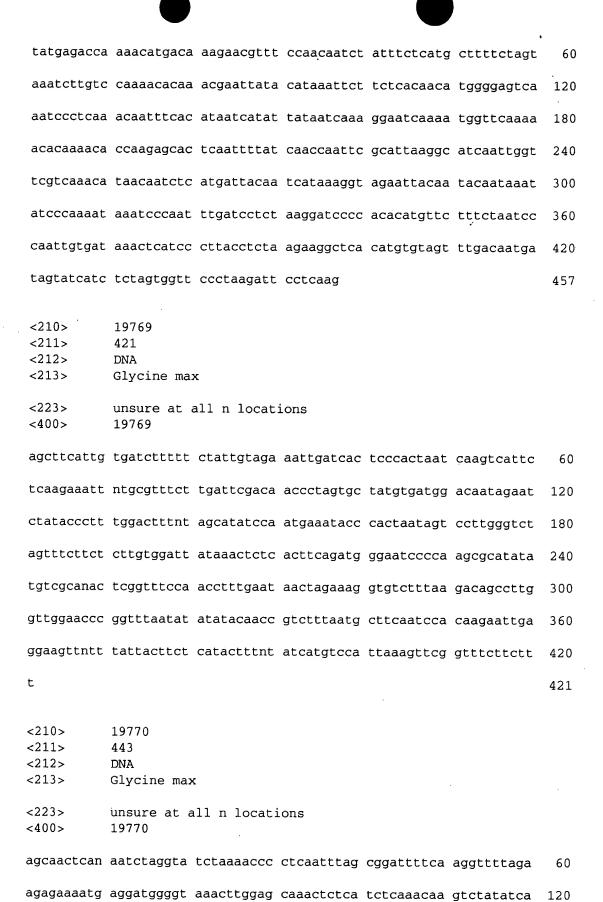
| ataaattago | c cgaatctaaa | atcttagaag | acgaaggagg | ggggggaaga | tcaaaactct | 240 |
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| ga | | | | | | 242 |
| <210> <211> <212> <213> | 19757 409 DNA Glycine max | | | | | |
| <400> | 19757 | | | | | |
| agcttgttta | tcccttatga q | gttgaatcgc | acatagaatc | gtttttttgg | taggctcgga | 60 |
| gtaaactctt | tacaatcata (| gttttgttac | tgtgccgacg | attctactcc | acaccataga | 120 |
| caacaccgta | atagtatgtc a | acgacagtgg | ctgggagagg | cggcatggcg | aagtccgtga | 180 |
| acgagggttt | cactagtgag g | gttggtcgaa | aatggcggac | atgcaacaga | aaaggatgca | 240 |
| tgaagaagag | aaagaatata t | ttatatcat | tgtggcaaaa | ttagttctat | gatgtggcgt | 300 |
| gccagcgacg | atggatgacg g | gtggcactga | ttatgtgaag | cttcatgcta | agcatagtgt | 360 |
| cgcaatcact | attggaacga g | ggcgtcgtg | tgagataaaa | atgattatg | | 409 |
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| tcttcttcca | tggcttattc c | ctagtggat | ggcgcctctt | ctcacctctt | ctcctttgtc | 60 |
| ttccgctgca | tctccatggt g | gaaaaccac | cattaaagga | cctcattgaa | actcaaagat | 120 |
| ccagcctcca | tagaagcccc a | caagcaagc | ttccatcaag | tggtaatcag | agcacaagag | 180 |
| cttcaagtag | gtgctcctta n | acctccatt | aattntttt | gctttacctt | ctcttccatt | 240 |
| nttgtttctt | cattnttctg c | atgtatctc | ctcacatgtc | ttgtgataaa | tgttgttaac | 300 |
| atgattcttt | agagtttcca c | tgattaaac | atgctataga | agctagattt | gattttctat | 360 |
| ggntcaaatt | tcttgttctt t | ttcttgaac (| catgaattgt | gttgac | | 406 |
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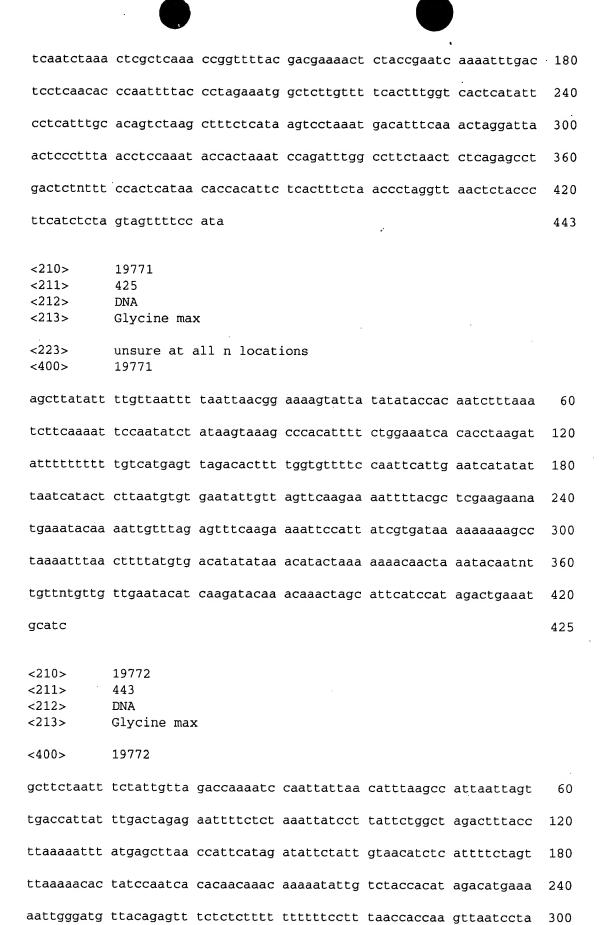
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| <400> | 19759 | |
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| ccttatgca | c ttctctctct ctcgaatttg ctgaggaaaa ttatttccgt gaagaaaatc | 120 |
| caagccgag | g cgcttccgta acgtttccgt gagtaattac gcgaagattc tcgaccgttc | 180 |
| ttcaagatco | c atcgttcgtt cttcgttttc ttcagtctac aacgggtaag tacctcaaac | 240 |
| caagcttttc | aattegttet atgtaceegt ggtggteeae attatgttge atgtatttt | 300 |
| attcttgtgc | tcgtcaactt tttatacccc ttgtgacgtg cttaagccat ttatttaagt | 360 |
| catttctcgc | ttaatctacc aataaaataa atttccatcg atc | 403 |
| <210> <211> <212> <213> | 19760 444 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19760 | |
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| agagggaagc | tccccaagtt ccaactccga acacggctcg accggccggt aattccaaca | 120 |
| • | cttccctcca aggccattgc cggaattcac cccgctcccg atgacgtacg | 180 |
| | accatecete ategecaate atttggeegt ggtaacteee ggaagggtee | 240 |
| | tttcccgaag tggtatgacc ctaatgcaac tngcaagtac catgggggtg | 300 |
| | ttccgtcgaa aaatgcttgg cccttaaata caaggtccaa catctaatgg | 360 |
| atgccggatg | gctgactnct caagaggatc ggcccaatgt gaggaccaac ccgctcgcca | 420 |
| atcatggagg | gggagcagtt aatg | 444 |
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| <223> <400> | unsure at all n locations 19761 | |
| ttagcttaag | taatgccagg gatacaccac ttccctttta gggtagttag tttgggatat | 60 |











| taatccccca | tttctctctt | taaccaccaa | gcaaacctta | a taactcctaa | ı tttgagggat | 360 |
|---------------------------|------------------------------------|-------------|------------|--------------|--------------|-----|
| caatttttct | ttgtatttat | tttaaggatt | aactttctat | ctattacaac | ttgagagact | 420 |
| aaattgagga | ataatgttca | acc | | | | 443 |
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| tgagaaccat | aggccggtgt | ctatgggtcc | tttgatatta | tgaacaatcc | actttgcatc | 60 |
| cttgagagga | gtactcatta | cagtctccat | gtatcaactg | atgactccag | cagcatataa | 120 |
| aacgtctagt | cttgtgcaca | tcatatcgta | aactacccac | caaactcttg | aactttgtat | 180 |
| catccacctt | tcctgcttga | tcgaactttg | ataacttatt | tctgcactca | atcggtgatc | 240 |
| caattggctt | gcaagcatcc | atcttgaatc | tattgagcat | ctggtgtgtg | acatagccct | 300 |
| ttcacactta | cgagatggca | aagtggctct | gtgagtgggc | gatgacgatg | gtgagagctc | 360 |
| ttgctgaaca | tgttgaggtg | tcttatcttc | ttcaagt | | | 397 |
| | 19774 421 DNA Glycine max | 11 - 1 | | | | |
| | unsure at al 19774 | II n locati | ons. | | | |
| agcttcttgt | tgatcaagtt g | gatccgcaag | ctagttacag | atgaagttga | tctgtataaa | 60 |
| tcaacttcat | ccgtaaccag o | cttatggatc | atctacgtca | actacggatc | aaatatagct | 120 |
| tctgcgggtc | aacaaaaacc t | atgtgaatc | atgccatagc | atacgcggat | caagctgaat | 180 |
| gagccgggtg | cacaaaaata t | tttaaaaat | acacggnggt | atttttgtct | tttcatgtag | 240 |
| ggtgttaggt | gcaccagcaa t | aatgctggg | tgctcctagc | aacacccttc | atcaaaatat | 300 |
| aacatccact | ttacaaagtt t | gtaagcctt | ttcagggcta | cggttccctt | gttacccgtt | 360 |
| tacaacacca | aattcatatc a | atatgctaa | tcttgtacca | aaatgtttgt | ctctacttat | 420 |
| t | | | | | | 421 |

| <210> <211> <212> <213> | 19775 433 DNA | |
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| ctattttgaa | a ttctttagtt cctgaatgta caaccttcaa attgttactc gttcccgtat | 120 |
| ttgttttctç | g caaaaaagaa aattaatctg aaacaattca ggctgaattg ttatcgttat | 180 |
| tattactcga | a accataagga ataacagcta aacaagtaat ttaaaatgta acttttaaat | 240 |
| tatgtggtat | ttttttaatt acaattttac ttcaatatct aattttgtta atctacttag | 300 |
| gtcgttgttt | aaatataaat atgaatttaa aggtgatcta ctgataatat aaagtacttg | 360 |
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| aaatgtataa | taa | 433 |
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| <211> | 420 | |
| <212> | DNA | |
| <213> | Glycine max | |
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| ggcgggttgc | agcaccggct ccgcttccct aactgtactg gaggcggttg cggtggcttt | 120 |
| atcctctatg | gttttctgga gttttaacat gacctccgag atggaagcca ttttatcttt | 180 |
| caaggccgat | agateggeet teatetgtte etgeaegeee tetteattat ecattttet | 240 |
| agatcgagtg | ttataggggt gccttggtgt tttcttagtt atgatgaaat tcctaaagaa | 300 |
| ataaacaacg | gtgagtatgc caccaaaaca tgagtatgca aatggatgat cggagcactt | 360 |
| ggatccaccc | caagattntt agataacgta atgagtccag aactttctca ttntataaaa | 420 |
| <210> | 19777 | |
| <211> | 434 | |
| <212> | DNA | |
| <213> | Clucine may | |

| <223> <400> | unsure at 19777 | all n locat | ions | | | |
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| ntgtacggcc | ttaagcaggc | ccccaaacaa | tggtttgaat | tactgcaatc | tactatcttc | 60 |
| aagccctttg | tgatgcaaac | tgggcatcaa | atgttgatca | ctgaaggtga | atttcaagtt | 120 |
| ttgccatata | tttgggccct | tatcatatat | cttggtggac | ctgcaagatc | aagtgcaaag | 180 |
| gcaaaatatc | gtagtttggc | ataaactact | atagaattat | cctggattga | gaccatgttt | 240 |
| aatgagttgt | aagtttcctt | caacacactc | attgtattat | gtgacaacca | aagtgttgtt | 300 |
| gctcttgccc | actaaagtta | gttattgaca | actgttgaca | cctaactcaa | tagattggta | 360 |
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| caaaacatta | ttgtcgtttg | aattagctca | gagcttcaga | attcaatttc | gatcgtctcg | 120 |
| atatattacg | ggtctcaatc | agacatctga | gtaaaaaagt | tattatcgtt | cgaatttgct | 180 |
| gagagcttca | acattcaatt | tcgagcgtct | cgatgtttta | tgggacttaa | tcagacatcc | 240 |
| gagtaaaaag | ttattgccgt | ttgaatatgc | tgagagcttc | aacattcaat | ttcgagcatc | 300 |
| tcgatatatt | acgggactca | atcagacatc | cgagtaaaaa | gttatcgtcg | tttgaatttg | 360 |
| gtcagagctt | caacattcaa | tttggagcgt | atacatatat | t | | 401 |
| <210> <211> <212> <213> | 19779 379 DNA Glycine max | ĸ _ | | | | |
| <400> | 19779 | | | | | |
| tctggaagga | gatcaacttg | atgttctatg | cctcttgatt | gtggtagtcc | atgaggaatc | 60 |
| tccataggaa | agacatttct | aaattcctgc | aataagggtt | gaacactagg | agaaatagaa | 120 |
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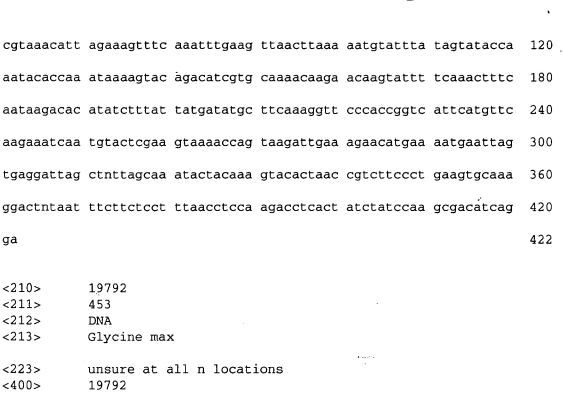
| agtggttcaț | gagcaggtaa | cattttcctc | acttcactcg | cctctgcaaa | ataattaaat | 240 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| tttctctcat | gtgtatcact | cttttcctcg | ggtgtatcac | tcttttcat | attccttttg | 300 |
| ggtgcctcac | tattatcttt | ctcttggtct | ctcttttctc | tcattctgat | ttggtcatca | 360 |
| cacacttctc | taggggata | | | | | 379 |
| <210> <211> <212> <213> | 19780 349 DNA Glycine max | κ | | | | |
| <223> <400> | unsure at a 19780 | all n locat: | ions | | | |
| agtttgcatt | atattgtgtt | tggnctgaat | ccttatgagc | aagcaatatc | aattattgcg | 60 |
| aaaagcttgg | cagcattcga | taaggataac | ttgatcccct | actttggatt | tggagatggt | 120 |
| atctctttaa | acaaatttaa | cttatggagt | gcattatttt | gaagttaaga | tatggctaaa | 180 |
| agaacttgca | gaaagaatct | gatttggaaa | tttgacataa | aaaatgtgat | taagatacaa | 240 |
| gttgcagacc | acatgtcttg | ccttgaaaac | aaagcactaa | agatgaacct | ttgcacattt | 300 |
| aggaaatact | tgctaatgag | caattacttg | gtataaaatg | acatgcatt | | 349 |
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| <223> <400> | unsure at a 19781 | all n locati | ions | | | |
| agcttgtatt | cattacttgt | tgagaaccat | aagccaaagt | cgattgttcc | tttgatatta | 60 |
| tgaagaattc | attttgcatc | cttgagatga | gtagtcatta | gagtctccat | gtatcaactg | 120 |
| atgagtccag | tagcatatag | aatgtctagt | cttgtgcaca | tcanatcgta | aactacccac | 180 |
| caaactcttg | aactttgtag | catccacctt. | tcctgcttcg | tcgaactttg | ataacttaat | 240 |
| tntgcactca | atcggtgttc | caattggctt | gcaagtatcc | atcttgaatt | tattgagcat | 300 |
| ctggtgtgtg | acatagccct | ttcacactta | ggagatggca | aagtgtcttt | gtgagtggtt | 360 |
| gatgttgatg | gngtgagctc | ttgctgaaca | tgttgaggtg | tcttatcttc | ttcaagt | 417 |

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| ntagtgcatg | taatccactt atgaatagac cccatcttcc tcataatata aaactagaaa | 60 |
| tcccacaaac | ataatcgtat accttctcaa agttaacttt aagcacaagg catgatttct | 120 |
| tccttctctt | agcatcatca accacctcat ttgcaatggc tacactatct aacatatttt | 180 |
| gctctcccat | aaaggcaatt tgcttttcac caataattnt agggagaacc actcttagtc | 240 |
| ttcttgctaa | aaccttagac aaaattttat acaagcaccc gattaaggat ataggcctaa | 300 |
| agttattaag | gccttgtgga tcatcctttt ttgagatgag aatgataaac gaaggattnt | 360 |
| ctcttcttgg | aattgctcca ttttcccaaa actcttgcaa catatttata aaatccacct | 420 |
| tcaatgtagc | ccaacatttc ttg | 443 |
| <210> <211> <212> <213> <223> <400> | 19783 381 DNA Glycine max unsure at all n locations 19783 | |
| | tatatggcct ccgtgataga agccatttga tcttttaagg tccataggtc | 60 |
| | | |
| | tgttcttgca ctccctcttc attatccatc tttctggatc gagtgttata | 120 |
| | tgcgcttttt tagttatggt gagttcccta aagaaacaaa caatggtgag | 180 |
| tataccacca | aaacatgaat atgctaatga atgatcagaa cacttggatc cacctcaagg | 240 |
| ccttttttag | ataacatgat gagtttcaga acttctcttt ttataaaaag gaacaaagct | 300 |
| tttatctagc | caagatcata caaaagtgtt acaacagaac gtaacggttt ctaattatat | 360 |
| gggccatcaa | atctatcatg t | 381 |
| <210> <211> <212> <213> <223> | 19784 420 DNA Glycine max unsure at all n locations | |
| | | |

| <400> | 19784 | | | | | |
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| tcttatccaa | ggctcatctt | nggggtgaag | ctccttcctt | catggcttat | tccttaatgg | 60 |
| atggcgcctc | ctctcacctc | ttttcctttg | tcttccgctg | catctccatg | gtggaaaatc | 120 |
| accataaaag | gaccccattg | aagctcaaag | atccagcctc | catagaagco | ccacaagcaa | 180 |
| gcttccatca | ttttcttctg | agggcaaggt | ttatggtaaa | ttgggatttt | ggctcaaggc | 240 |
| ttgtaacacg | gctggacatg | atatatgtca | gggtttggat | cggttcaagg | gtaaaagggg | 300 |
| atgtcccaca | ttatttccat | gacacaaatg | caacaatgat | gatntggaaa | ttttatacac | 360 |
| ctatgtggac | actcaagtgc | canactttta | tggtcatgtg | atgctatggc | tcaggattca | 420 |
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| tcgacgctcc | ctcgacggtg | gacggtggtt | cgtggctcct | cttcacggca | acaagaagaa | 120 |
| caagaatcaa | acgcaagctc | ctcttcacgg | caacaatggt | tcgtggctga | ggaagaagaa | 180 |
| gaagaaccaa | acgaggaaga | agaagaaggt | agcgcgagaa | agacaggtct | aagaggtcac | 240 |
| caacacaata | tgaaaatggc | tctg | | | | 264 |
| <210> <211> <212> <213> | 19786 392 DNA Glycine max | | | | | |
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| actgttcttc | cttcccgcga | tgcttctttt | catgtccgcc | tgagtgggct | tatagcctaa | 120 |
| accatacttc | ccacggtttc | cttgagcatt | tatcaggcta | gttatgccgc | cgttgtcttt | 180 |
| gcctaaaccc | atcccgggtt | cataaccgtt | ccccaacata | actcgggcca | tcattaccgc | 240 |
| tgcatcggac | agacaaggct | tgccaaagag | ggagtccaca | gaggaaatgc | tgaccacctc | 300 |
| aaaagactag | aaagcagtct | ctaacgattc | ttctgcggct | tccacataag | gcatggagga | 360 |

| | tggggagctt | accaagatgt cttcctcgcc tg | 392 |
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| | <223> <400> | unsure at all n locations 19787 | |
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| | aacattttcc | ctttttgcca ggaaggggcc ttgggctctt gacgtaagcc ccccccagg | 120 |
| | ctggactggt | tcgacccgac taaggaccag gattagtgag cttaccctta catcgttgga | 180 |
| | agacaacaat | tcaccgaggt agaaacgcac acagctatca cggatgacta acatgaagaa | 240 |
| | acacgccatc | taaagctcaa ataggaccac gcttctaaaa gactagaatt ctaccaaaac | 300 |
| | ttaaaggtgc | aaagacgacg gaaaaattct ggcagcaaaa taaaagcaca taaatacccc | 360 |
| | taagccacc | | 369 |
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| | <223> <400> | unsure at all n locations 19788 | |
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| (| ctatggcgga | tctatacatt cgctcttatg gcgagctttc ccatgcattt gcagcggaag | 120 |
| ç | gcgaaggatc | ataggtcatg gtagacacca ttcacagagg aataagccct ggaagatgga | 180 |
| Ć | gctttcaccc | ccagactgtg cctccggtaa gaaactcgac gaggaagcct taatggacga | 240 |
| ζ | aagacagat | ggaatggggg agtccgaatt tgatcgaata caccacggcg agaagtggaa | 300 |
| | ctcgaggtg | cgcctcataa gactcttatg catctaagtg accacaagcg tacccatgct | 360 |
| t | ctatttata | gactatgtac ctacttgaaa gcttctagaa atacttcctt gacaacttca | 420 |
| - | tgaaaaact | cettgaaage teagetaget accaeaceae ttaaagetag ceacetetta | 480 |
| - | aaccn | | 486 |
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| ۱ | 210> | 19789 | |

| <211> <212> <213> | 387 DNA Glycine max | |
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| agcttgttat | tctattttat ccaaacaact tcacaaatct cacatgcttt caagaaacaa | 60 |
| cctcttcaat | cttccatgtg taccttcaac cctttaaagc atgttaatgt aatgtcaaca | 120 |
| gaagattatt | cttataaaaa acaaaatgtg aaacaagtag gcaattacat gaatgtacct | 180 |
| attagttgtt | gtgtttccaa tatgcataac ccaatcagtc catgcttcta caaacttctc | 240 |
| | atcaacaatg tatagtaaac atactaaaaa aatactaaaa actatgaaca | 300 |
| | ccctaatttc gtccggagac cattatttgt tggcatgcgg ccttcatttg | 360 |
| actacctcaa | aatgtttaac acccatc | 387 |
| <211> <212> <213> | 19790 439 DNA Glycine max | |
| | | |
| | tttatttcca cttttaagca acatattagt accaagcaat gctatcaaac | 60 |
| | aaaatacaag agcaaaaata ttacacaatt agcattttga atgtacaaaa | 120 |
| | aataacaagt aacagggagc agtgcacttg agaagcaact acatttcacg | 180 |
| | gtatttgttt tatgcagttt taaacaaaat cccattttaa aatcattcat | 240 |
| | ttctcaaatc ttattcatcg cctaacttct gaaataaata tcaaataaca | 300 |
| gggaacaaga (| actttgataa gtcacaaagc tgattgcatc atactcaatt tgagagttct | 360 |
| caccaaattc a | agcggcaagc aagttattcc attggctcta gagtcgtcta gccttgatct | 420 |
| ttggcacgta (| Ccatataat | 439 |
| <211> 4 <212> 1 <213> 0 <223> 0 | 19791 122 DNA Glycine max Unsure at all n locations | |
| _ | caacctgca aatttcaaga caaacaagag caatgcaaaa attcacaaat | 60 |



tatcattcta tctctcanaa agtatgaaag gaagttcaag tgatgcaggt aattcatacc 60 cctagtagac aaatatttct atcttacctt catgtagaca tctgtatctg gatcaggtgt 120 aatatttgct tcttttctc ttctgaacac ctctgctagc aaatctgcag aatgtgaaat 180 aattttagtt aaaacttgat ttttcatatg aattttggct cgatttccaa attgcagctc 240 tgtgggctaa ccataaggag gcccaactcc ttggactcat gctgagaagg ccagggtttc 300 tctgactgtc atttccaa tatgaagatc attttgactt acaaactcat tcatcccatg 360 accattataa gtcacctttc cagtgaactg atcaagaagc caaacctaat tagctacaaa 420 tccaataatg gaaaatatga gcacaaacat cat 453

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|-------------|---------------------------|
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| <213> | Glycine max |
| <223> | unsure at all n locations |
| | |

10702

19793

-210-

<400>

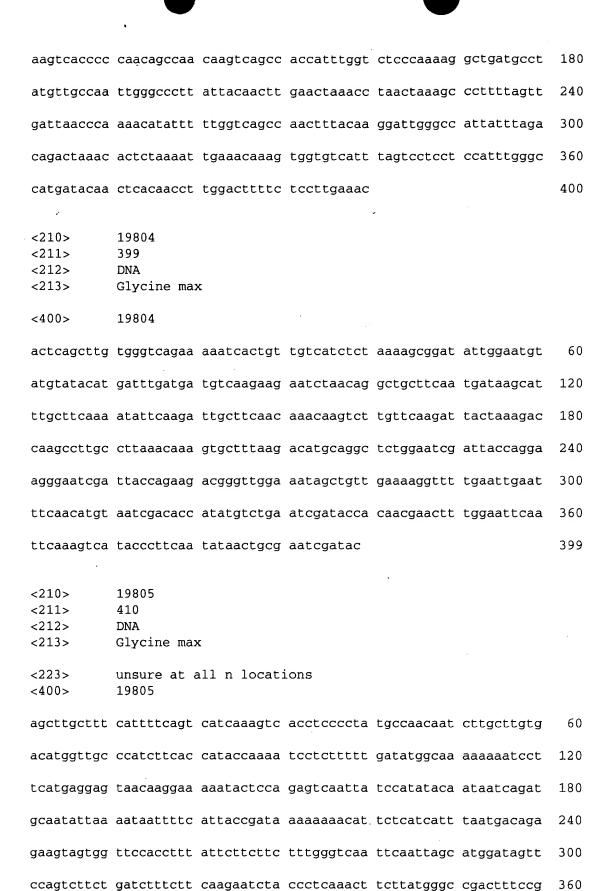
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| gatcttcta | ac atcctaggtc ttcccgttcc ttcatctggc ttatgttctt catgtagcat | 240 |
|----------------------------------|---|-----|
| tcagactga | aa tgactctatg aaattacgtc gctacttcca catggtacgg gtaacgtatg | 300 |
| agacatctc | ct attiticccg gngggaatcc ttagaattac cacagcttag cinicaatto | 360 |
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| tcaagcttg | t gaatttagtt ttgatgcagc aagtgaggga aaacaattct taattgatgt | 60 |
| ctcaaaaaaa | a aaagcagcgg attcagtagc acgggctgca ataagagctc ggtgtcatta | 120 |
| tgttaataaa | a aagtggctcg gcggtatgtt aacgaatcgg tatactacag aaacacgact | 180 |
| tcaaaagtto | c agggacttga gaatgcaaca aaagacgggg agactcaata gtttttcaaa | 240 |
| aagagatgco | c gctatattga agagacattt agctcatttg gaaacatatc ttggcggcat | 300 |
| taaatatato | g acggggttac ctgatattga ataatcgtcg atcaacaaga agaatatacg | 360 |
| gctcttcgag | g aatgtataac tttggaaatt ccaacaattc gttcaatcga tacaaattgt | 420 |
| gacccggacc | c togoogatat ttoaattooa gotaatgatg a | 461 |
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| agctttgctt | aatgagaatc tgttcttcca acaggctgtg caatttggca tctgaagtct | 60 |
| gggagctctg | ggcagatggg tctgctaaag ctagaagtgg ggctgaagta gaagatgcaa | 120 |
| ggatgccagc | tattggtgca aaggaaaagg gaacatcagc tgctctgagc ttgctcttcc | 180 |
| ttgcctctgg | aaaattaact gtttggtcat tcgcattcca acagttcttt atgatatacg | 240 |
| ctaagtcaat | gaccggtctt aggttcttgt aggaggtaag agcatcagat ccaactcccc | 300 |
| tcaatctaca | caaggctgtg attaaagctg ggaagcctaa gcgagaagag ttagactgag | 360 |
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|---|----------------------------------|------------------------------------|--------------|------------|--------------|--------------|-----|
| | <223> <400> | unsure at 19796 | all n locat | cions | | | |
| | atgcacggaa | . aatgtaatta | a tganattgag | atgeçegttt | attcaccatt | tcctagttaa | 60 |
| | ccttgcatta | agtaccatgt | tcaattattt | tgtttctaag | , tgaaacgggt | ttatgatccc | 120 |
| | aacatggttg | gctcgtgggg | cctaacacat | gatacttaga | ı atgtagtgtç | g aagtttcacg | 180 |
| | ctttcccctt | tttgtgtttg | ttttgtagaa | ggaaaacgca | aggatgagca | a aacattgaaa | 240 |
| | caaatggtat | ccaattttgc | agatacaaaa | gtttg | | | 275 |
| | <210> <211> <212> <213> | 19797 416 DNA Glycine ma | x | | | | |
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| | agctttgttg | tggaaggaag | atggaagaaa | gaagaggttc | gaggagattg | gtaaagagga | 60 |
| | tatacatttg | aactaaggct | tgccacatgg | tttgctgaca | tggactcaga | ttttccatgt. | 120 |
| | tatggactaa | gacttgccac | atagacattt | gactaagagg | aaattagatt | ttaaccgtag | 180 |
| | ggacttattt | gcataacaaa | tgtaaagatg | aggattattt | ttttcatttc | aataggatag | 240 |
| | agactaatat | acaaaatgac | tacaaagata | gggaccaaaa | tgcctattta | ctaaaaataa | 300 |
| | aagagaacaa | tgtcatgaac | tttctaaaac | tcttgttctc | tccttatctt | gtaccaaagc | 360 |
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| | <223> <400> | unsure at a 19798 | all n locati | ons. | | | |
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| ç | ggaggcaacc | catctctttn | tattttcatt | aatcattgca | tatagttagg | attaacttat | 120 |

| ttgtgaattg | ttgagtaagt | catcagcatg | tttagttttg | aaaatggggt | tgttaaagct | 180 |
|-------------------------------------|--|-------------------|------------|------------|------------|-----|
| ctttgaagct | gtggatgaaa | aataacttag | aaaatttttc | agtcatccac | tcgctcagcg | 240 |
| cgccctgtgc | gctaagcgaa | ttatcattca | tgcgctgagc | gagtctcaac | tcgcgctaag | 300 |
| tggatcaacc | cctacctatt | agctgatggg | gtctcgctaa | gcgagacctg | tgcgctaagc | 360 |
| ccaaaaacct | ctctgaaatt | gcatttaatg | gaattagggt | aagcaagtct | tctttctaag | 420 |
| cgcacaacat | tgtctcgct | , | | | | 439 |
| <210> <211> <212> <213> <223> <400> | 19799 87 DNA Glycine max unsure at a | c all n locat: | ions | | | |
| | | gaatgaagct | ctgataccac | ttgttagaca | agtggcctca | 60 |
| | gaagggggg | | | | <u> </u> | 87 |
| | 323333333 | 333333 | | | | 0. |
| <210> <211> <212> <213> | 19800 442 DNA Glycine max | s | | | | |
| <223> <400> | unsure at a | all n locati | ions | | | |
| tctanccttt | tccttccttt | ctaccacaaa | ggtggagtta | ttccacatac | ataaaaggcc | 60 |
| accagcagct | tccacagatg | gaacaaaatc | ccaatgacca | gtggagtctc | ccgaaatggc | 120 |
| ctggcaaata | cttttattaa | agttctccct | cttggtttct | tggaggcaga | caagatgcac | 180 |
| tttgtgctta | caatgagcct | tctaacagca | gcccacttga | ctcccctccc | ctaacctcta | 240 |
| gaattatagg | agagaattat | cataattgct | gagatttaat | tcccttttct | gttgccatca | 300 |
| aatcatcttt | attctccata | tccagcagta | gccccttaac | cttgctatct | tcttccttat | 360 |
| aagacaagcc | catttccttc | aagatgtcac | attgcagctg | tatagggtcc | tcgtataaag | 420 |
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|----------------------------------|--|-----|
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| cactgcgccc | c ccaaatacgc aagtaagaag agataatatt ccgggctctc gggtctgtga | 180 |
| aatgcattca | tatcatgcat cgcataaaca tctgttcatg gcatcataat gaacatatcg | 240 |
| atgctgcatt | tgtctgttat catattacag cctcacattc tgcatgagtc atggcatcat | 300 |
| catgcatatg | g cgttcaacac actttttgat ctgcacaatt ggataccatt tggtttcatg | 360 |
| ttagctca | | 368 |
| <210> <211> <212> <213> | 19802 393 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19802 | |
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| gagttgtgct | ggaagtttct ggagaaagag agagaagatt tggcttttaa aatggttttt | 120 |
| cttttcttt | ttcattntct tttttaaaag caattccaca tgtcattttt ttaattggag | 180 |
| caaaaagggc | c ccacctttac ctttgacttg accaaatact cagccataaa agaagaaaaa : | 240 |
| aatgggacct | ttttggatgc tgaaatcttg cctcggtttg cgtgccgcct cttcggttcc | 300 |
| agttcttcgt | gtttetetge accegtegag geceatttte aaggtaggea atatatatat | 360 |
| atatcaaaat | gctcagaatg agaccctgag cat | 393 |
| <210> <211> <212> <213> | 19803 400 DNA Glycine max | |
| | | |
| <223> <400> | unsure at all n locations 19803 | |
| <400> | unsure at all n locations | 60 |



| | aactaaagcc tttgng | gtga gacttggato | : tttcttgtga | | 410 |
|--|---|---|---|---|--------------------------|
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| <223> <400> | unsure at all n 1 19806 | ocations | | | |
| tgcttctaca | actctattta taatat | ttgc tgcaatcttt | atttcctttc | ccaagagtga | 60 |
| ctatagtaag | ttagaatgac aaatca | tgct tcttaccata | tccttaagag | ttttgtttca | 120 |
| tctttcagct | actacateca tgetag | gtga ccacgacatg | gtgtactatg | ggacgatttc | 180 |
| acattcctct | aggtacctag caaaag | gccc cggacgttgt | tcacttgaac | cgtcatatct | 240 |
| gccatagtat | tcaccaccac ggtcat | atct aacactcttg | attcttttgt | tgagttgatt | 300 |
| ttcaacttca | actntaaatg ttttga | acac atccagagat | tgttatattt | catgtataag | 360 |
| aaacaagtat | gcatatctgg agtaat | tgtc tatgaatgat | ataaaatatt | gttgaccatt | 420 |
| ccatg | | | | · | 425 |
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| <210> <211> <212> <213> | 19807 369 DNA Glycine max | | | | |
| <211> <212> | 369 DNA | ocations | | | |
| <211> <212> <213> <223> <400> | 369 DNA Glycine max unsure at all n lo | | tcttacttat | aatcggcctc | 60 |
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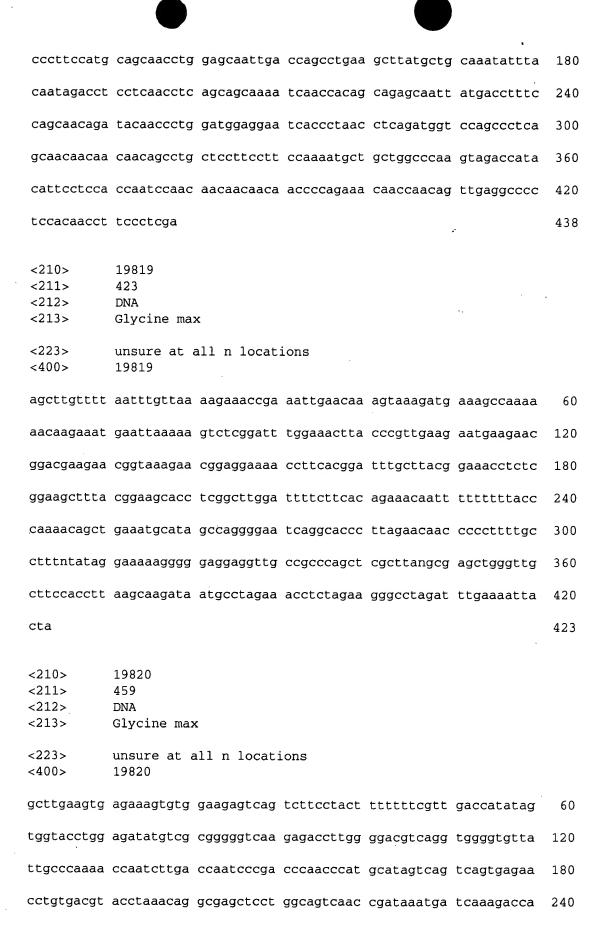
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| gagcgttgca | aggctaaaga | aggacgagat | ccagaggctg | aagaaggaga | tcaattagct | 180 |
| ccgacgtccg | gcgacagagc | tgcatgactc | agagacaagc | gcgacgctga | agaacctcct | 240 |
| cgaagaggga | gaaagaatgg | tgacattcct | agagacgagc | gcgccagcac | caccatcacc | 300 |
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| ggttcacatg | tgaattattg | tgaattgagc | tttgttgctg | agacaatgtc | atcaaagctc | 180 |
| tatattgttg | taaagccaat | ctcattatac | tattatgtgt | atcttgactt | tgtactggaa | 240 |
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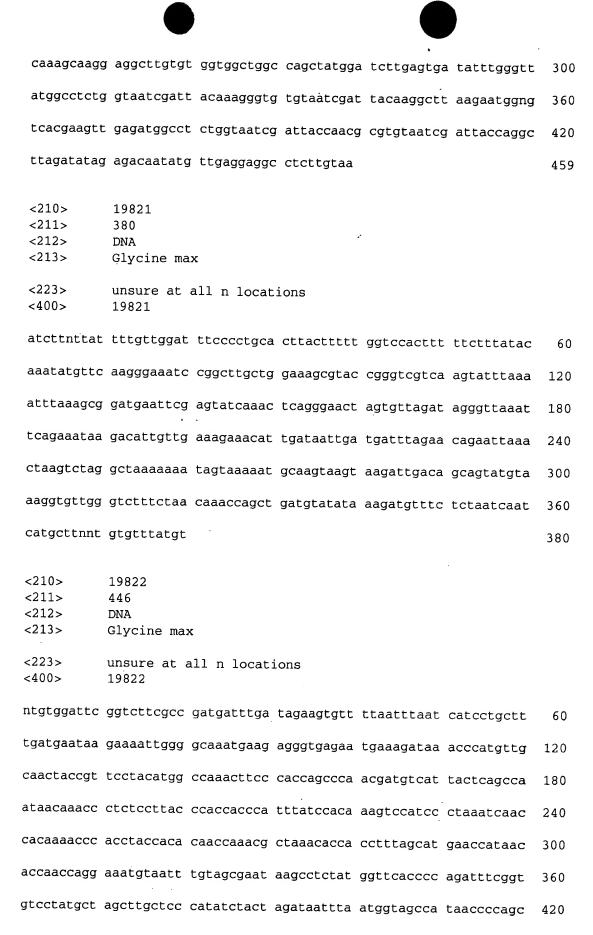
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| gataaaaatt | ctttcatctt | tttattataa | aaaaaataca | atcatatgta | tgcctataca | 240 |
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| acatttgtgg | gtggccaaat | attacacaac | tcatggatgg | ctgctagtgt | tgacaatcct | 360 |
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| attcttg | | | | | | 427 |
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| acaaagaagt | atgtattata | attttaatgt | acataccaag | tgaatggtgt | gatcctccaa | 120 |
| gctaatagaa | ccacctgtgt | gcatacccac | cctttttaga | tgcccttttt | tttgtctatg | 180 |
| cacacttgga | acgataccca | ggtgcattcc | aatgtaataa | tagattgttc | cagacacgat | 240 |
| ctccaatcaa | atatgacctt | tgattttcat | ttcgggcatc | cataaacatc | ttagagagtc | 300 |

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| ctgggtttt | tgggctcact tcctttg | ıtgg agtacagacc | caaaatgaga | tgacccatat | 120 |
| gtctcattta | tttaagtgga gagagg | ctat attagagagt | gagatacagt | gagagagact | 180 |
| catttgagag | ggaaaaaatt tacaaa | cat tgagagagat | agaatgagag | agaatatcat | 240 |
| tgtgatattc | gcatacccac tagagag | gegt ttttcagatt | gcaacttcgg | atttgttcac | 300 |
| cattggatcg | ggctgatttg aggacaa | actg gttctacgtg | cgtgatactt | caaattatct | 360 |
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| | aactgatgca | ttggtaactt | ggaacccttt | tgccttgaat | ctgaaatctg | tcctggcgca | 60 |
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| | gcctgctcct | tccttccaaa | atgctgctgg | cccaagcaga | ccatacattc | ctccaccaat | 360 |
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| | | cggtcgaaac | | | | | 120 |
| | agaagcgcct | cggcttggat | tttcttcacg | gaaacaattt | ttccaagcaa | attcgaaaga | 180 |
| | gagagaagtg | cctaagggac | cgaacccttt | ttcacttcac | ttctccccct | atttatagaa | 240 |
| | aattggggga | gaagcttgcc | acccagctcg | cccaggcgag | cagggttgct | tcctccagaa | 300 |
| | gcaacagcct | tctagaggaa | tcttctggag | ggcccaagtg | ggcctggttg | ctatttgcac | 360 |
| | ccccattttt | actaagtaca | cccctggcc | ttttttggtg | attcttttnt | cgtaaagtta | 420 |
| | cg | | | | | | 422 |
| | | 19818 438 DNA Glycine max 19818 | | | | | |
| 1 | tgaaggcaaa | ctggatgcgt | tggtcaactt | ggtaacccat | ctggccttga | atcagaaatc | 60 |
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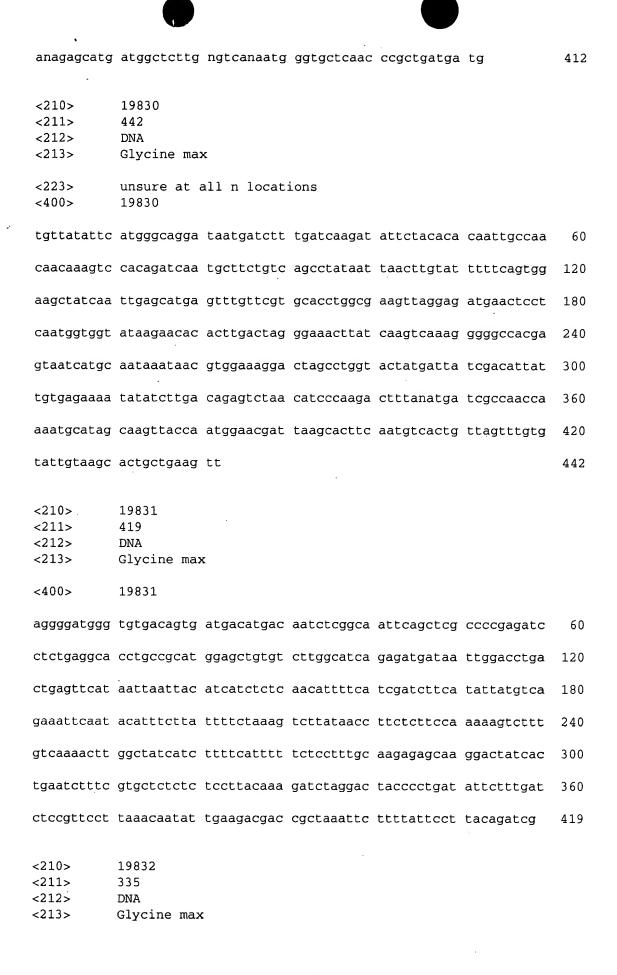




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| tgttatttga | ttttaagatg aaatctaaca tggtataaag ccttacaacc catctttgtg | 180 |
| tattgcctat | ttttctacat tggtggagtg gggatgttga aaagtcgcac atcgtttgcc | 240 |
| ttaattctga | gggtgcaact tatatactta ttgggaaact ttacttaaga ctaatcgatg | 300 |
| ttatgaatct | tatgatgaaa cctaacaact tgcaacgaag agacttgtcg gttgctactt | 360 |
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| cggtttcaat | ttcaattgga gtgtccgatt gtgattgtgt taatgtgatg gattttggtg | 180 |
| tggtgagtgt | ggattntggt ttcaggtacg ttgggatctg anggagggtt gtgttgttgt | 240 |
| gttgtgttgc | gtgtttgtga aatggaggac atacgatacg ggaaaggcgg gctgtgaaat | 300 |
| gagatcggga | tcttccaccc ggaacaatcc aacaccggag tctagtgctg ggtggggctt | 360 |
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| tntgcgttta | attactcat | 439 |
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| agacggtgat | gaaactgatg atatccatga aaatgatcct attagagaat tagacatggt | 180 |
| tcttgacgta | gctcctatgt ggagcttgta ggccttggat attcttcatc aatggagtcc | 240 |
| tttgcttctt | gaagatgaat gacagcagaa tggagaagga agatgattgg agatgccact | 300 |
| tcaaggagaa | gatgaatcaa gaagaagctc accaccatag gaagccatgg ataagagctt | 360 |
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| aaaagctatt | gatctttctt attcctcatc cagagaagct gttgctctct ctgattcccc | 180 |
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| ctatattagt | ttcagtacat tattttgagt aattctggtt tctgattata attatgtact | 300 |
| tatactttct | gtgattagcc actatgcttg gtttgaagca tatggaacaa tttgacttga | 360 |
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| cattacctgc | caaattggca attcctatgc agacatttgc aatgtctgat ggcactccag | 180 |

| cacttttaaa | aacagttgaa | gagaaataaa | acacagcatt | tataccagat | agctgttgta | 240 |
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| aagcaaatag | ggttgatcca | ataaaaacaa | ctgcaaaacg | aacttgtgaa | taaaatataa | 300 |
| cttttggaaa | cctaaggaaa | ccagtgacca | gataacaaca | atggcaaata | cactcacaag | 360 |
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| agattgaata | actatctgat | gattcaccct | gtaccttcga | taatttttca | ctctgctgct | 240 |
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| a | | | | | | 421 |
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| cacaccttgt | cactctcttt | cttcgccatt | ttcaaattct | tcttcattct | gaaccttatg | 300 |
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| gacccacac | a gtatcatgc | c cactactagt | cttgggtaaa | a ctagatacaa | ı aatccataga | 180 |
| tatgctctc | c catttgcatt | t ccggaatctt | caatggctgd | aattctcacg | atgggcgctg | 240 |
| gagctaacc | t aagcctttga | a catgtcaaca | tcttgctatt | atteggeaca | tcttattcat | 300 |
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| ttgataaagt | caaaaaccat | ttgaagagtt | atatattttg | atttattcag | aaataatcac | 180 |
| tggtaatcga | ataccatatc | agtgtaatcg | attatcacaa | agctcttatg | taaaacaatg | 240 |
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| acccccccc | a acaacaggaa ccggagcaca caccacacaa cccaacaaca cagagagcag | 180 |
| cgaccgcac | gaacaccacc cacgccaaaa gaccaacaca gacgccacac gaacaaaca | 240 |
| gggcggccca | a acgccaccga agcacaacaa gcgaaaaaaa ccccaaccac aaaaaaacac | 300 |
| gcgagacaco | agagacgaac ccaacacacg cgcgccagcg cacccgacac accaaaacgc | 360 |
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| agggattaag | agaccgaggg tetettgttg tgaaagaatt etaaacacaa aggaagggtt | 360 |
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| ctaagctcac | ctccttgaga | tgagaagcta | gagcttagct | acacacccc | tataatagct | 120 |
| aagctcaccc | ccatgacaaa | aaaaagatga | aaatacaaaa | gaaaagtcct | tactacagag | 180 |
| actactcaaa | atgccccgaa | atacaaggct | aaaaccctat | actactagaa | tggccaatat | 240 |
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| ttagcccatt | ggctcanaat | ataccctaag | gctcatgaga | accctagggc | cttcccttgg | 360 |
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| cattgatcca | gtcaagcttg | atgagttgga | aaatgaggcc | gcaaatatac | tgtgccagtt | 180 |
| ggagatgtat | tttccccccg | ctttctttga | catcatgatt | cacttgattg | tgcatctggt | 240 |
| cagagaaatc | aaatgttgag | gtcctggttå | tctactgtgg | atgtacccgg | ttgagcgata | 300 |
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| tatgtagtt | g gcgataatgt | acaatccttt | aaatagttta | tctttctgtc | atgcagtcgt | 300 |
| gagtttacc | atcgtgtgaa | gtctgtatct | atggcaatgt | tcacgtcaca | agaagttgat | 360 |
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| cctcactact | ttatcaactc t | ttccccagt | tgcatccgtg | tgtgtatata | tatatatata | 180 |
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| gtgcgaaaaa | agagacagaa a | agaaacttat | tcttttatta | atataaacaa | tctatatctt | 360 |
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| ctatttcaat | ttcaatgcaa g | ttgcaagtt | cccttaaaaa | tgaactttta | aataatgatt | 180 |
| caaatagaac | aatctgaata t | aaatataaa | tcaataataa | ataaacaagt | ttaagggaag | 240 |
| aaaaagtgca | aactcagatt t | atactggtt (| cggccacacc | cttgtgccta | cgtccagtcc | 300 |
| ccaagcaacc | agcttgaaag t | tccactatc (| ttgtaaaatc | cttttacaag | ttctgaacac | 360 |
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| aattaagtgt | tgagaggact | ataatttatc | gatatcgtgg | tccattctta | tccgatattt | 180 |
| tgactgcaac | ggcattgtct | gaacctaatc | caggaaccaa | agactagaga | taatatttta | 240 |
| tatacgtgac | ttctgggtct | aaaaatgttc | aatttgtgtt | atgttaatag | cgtgtaccat | 300 |
| ctatgcaaag | agcgtgtttc | ttgagtgcct | atatatectt | tagatagtat | astaattatt | 260 |

| ccgtttaga | g tggctttcac catctgtgaa taaaactatg caactgtagt accatctact | 420 |
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| gaacaccttg | gctgtatcaa aggactttca caacctttgt gtgttgccct cgctggaaag | 180 |
| agtgattett | tccttccttt catcatcacc cttgttcttt caaaccacaa ttccagaaaa | 240 |
| tccacctctg , | cccagaatta tctcgtggcc ataactccca ttntacgcac tcaaattaag | 300 |
| ttattcttga | gcctaaattg aatttaaaac gagacctttc acctcgtttt ggaatcacct | 360 |
| tatttggagc | cctgtncttc agttattgcc attctatatt tct | 403 |
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| | ctagagtete tataggegag geacegeeta tgaeagtteg aettgagaea | 180 |
| | aagagtetta caaaccatca gggteetatt eteageeact geacataata | 240 |
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| gcatgaatag | ttcaaagctt ctctatatag agagagcccg acttggacat tcttgagaca | 360 |
| agatatgacc | atatagaccg ctgggcgctt gcgttgaaca attccaaccc cacggcatgt | 420 |
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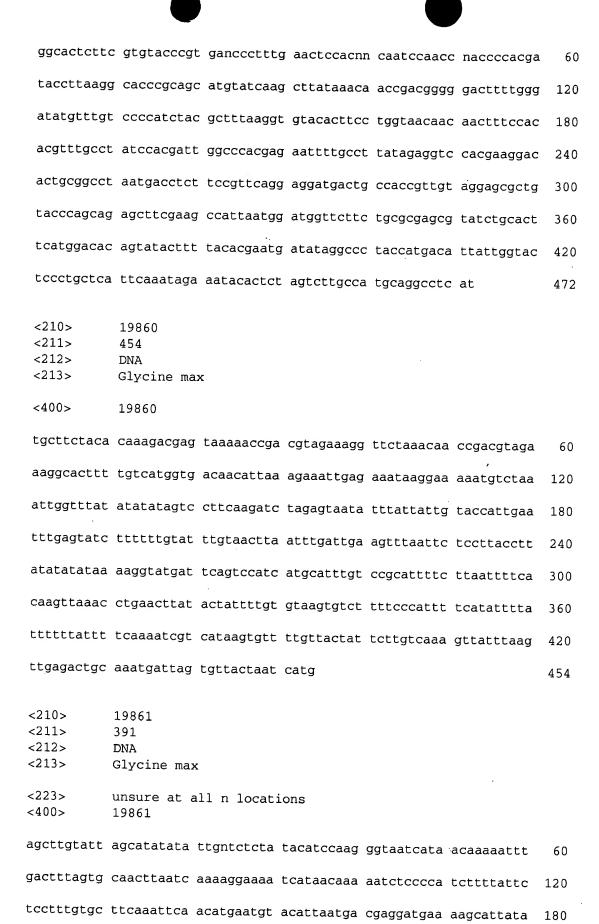
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| tatcatggt | t aaaagtgcat aatatatttt cctggtagca attaattaag gtgcaagtac | 180 |
| aatttaaaa | t gcacaacagt catgtagtaa ttcatattta gaatatgtaa taattcatat | 240 |
| atcctacat | t cataagacaa aatatagtaa gtttttatga accatagtta ggcatcctac | 300 |
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| ttgcccttct | gtgcaacaat ctgaagcaat tgaacagcct gaagcttatg ctgcatacat | 180 |
| ctacaatata | cctcctcaac cacagcagca aaatcagcca caatgaaaca attatgacct | 240 |
| ctccagcaac | aggtacaatc ccgggtggag gaatcatccc aaccttagat ggtcgaatcc | 300 |
| ttcacaacag | tagcaacaag aaccttattt tcaaaatgtt gcttgtagaa gcaaagcttc | 360 |
| atgatgaatc | cagattgatt caaagatgtt ctgatgataa caaagatgaa tgacaaaagc | 420 |
| tcaaggtcaa | tcaaagaatg ag | 442 |
| | 19849 382 DNA Glycine max | |

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| ctaccacagg | g agggagaett agceataeag eeeteaaage eteaetettt tteeaetea | t 120 |
| aacaccacat | teteaettte caaceetagg ttaaetetae ettteatete taatagttt | 180 |
| tccatgagca | a acttcagcat ataaacatca caaacatcat cacaaaaacc ctaaatagaa | a 240 |
| tgggtatgtc | taactcatcc agacatggca atttcaacaa gctttcaaca agagtcttca | a 300 |
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| catatcagca | acaagtgtct tttcagcttt agtcaatcga ccaacatatg gatgtccaac | 120 |
| taaagacttg | gccaattcat gaatatgaat gcaacacatc aactttacca tccaaccttg | 180 |
| tcctccaacc | actggtttgc cacaaagctt gaagggacac ccacatttcc taagtccagg | 240 |
| atctcttcta | acaaatgcag tcttcctaca cctatactcg | 280 |
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| | Glycine max | |
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| gcgcaaaagc | ttttgagtct ctaacttgga ttttcagctt gttacattct cctattaaca | 120 |
| catgatettg (| ctcttatgcc ttcttcaagg caacatcctt ctcctctatt ggctctaatg | 180 |
| ggttgatatt a | acctagggac tgcctcaatt ctagaaaaca ttttatatgc tagggtaaca | 240 |
| tgggtgagat a | actctatcac ccctaccaag attgatatgg tagaatctaa gctggtatga | 300 |
| ttgtttttta g | gaacaatate ecaattetea aacaaaataa gaegateaat eteeteacte | 360 |

| atagacato | a atatgagggg accgccaaac atcctggtac gaaccaagct agaatt | 416 |
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| ccaaaacca | a gettgaceaa teeegaceea accegggeat agteggteag tgagaacetg | 120 |
| tgatgtacc | t aaacaggega geteetggea gteaacagat aaaaggataa caagaccaca | 180 |
| aagcaaggaq | g gcttgtggtg gctggccagc tgtgaatttt gtgtaatatg tggatggtgg | 240 |
| cctctggtaa | a tcgattacca agggtgggta atcgattaca aggcttataa ttgatgacag | 300 |
| gaggctaaaa | a tggtctctgg taatcgatta | 330 |
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| ataatttgat | gctgatttgt gtgcaggtaa gttctcatgg cctgagttgg ttggggtgca | 120 |
| aggaacggta | gcggaggcta caattgagag ggagaatcct tcggtgaatg ctattattgt | 180 |
| | tccgtggtca caacggatct tcgaagtgac agggtttggg tttgggttaa | 240 |
| | attgttaata gagttccaaa aattggatag gaagtgttta cactanagca | 300 |
| | tatgttctaa taagttatga atatatagta atatttaaat aaggagttat | 360 |
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| | 19854 253 DNA Glycine max | |
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| cagcaacgt | t tgcccttttg ttgatgtgct atctatttta gttttagtat ctttctaaa | 180 |
| agttgggtg | go ctatatacag gtatattatg gaaatgtttt gaaaatccat tototagota | a 240 |
| caaagttgg | rg caa | 253 |
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| tgttttaag | g gtagcatttc ttggtaaaac taactttcca aatgtttgcc ttcgcaggaa | 120 |
| tggccccgaq | g gaagettgee teaaagaggt eeaagaagga eaaggeggee gaaggaaeta | 180 |
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| ttttgtgtct | ctcttctccc ttttccccaa gaacgaagga ctaacctcat gaattctttt | 180 |
| gtgtctccct | tetecetttt caaggaatte aaaaagacae agtetgagaa ttettttgat | 240 |
| tettecettt | cccttaaaca aaagatttca aaggactaat cgcctgagat atcntttgtt | 300 |
| tcccttttac | aaagtttcaa aggactaacc gcctgagaac tttgtcttaa cacattggag | 360 |
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| tgcaaagca | c agcaatgtgg gatcagtagc cccatcctta tagctaccat catcaatcat | 120 |
| gctctcccc | c aacaaagctg caattatata taaaagcttc aacgtaagat tgattttgac | 180 |
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| | t ggcacttaca actgaagatc cttgatcttg gagagcaggg atttagttgc | 300 |
| | g tgctttntaa acagcttcag tttctcttcg tcaagctttt gacaagaaac | 360 |
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| ctgcagcagt | gttggagtat ctatgattcc aagcctacaa tttcacaaac aatacatatt | 120 |
| cacaggtcca | aaaactgcag tgtcaagtaa actgaaacaa gacattcatt agctgtgtga | 180 |
| aggaatccaa | ctccatcttt aatgtaagat tacaacttac aagtggaaat aaatccatag | 240 |
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| | acctaacaat aggtatgagg catggacacc actaatttct caattgaaag | 360 |
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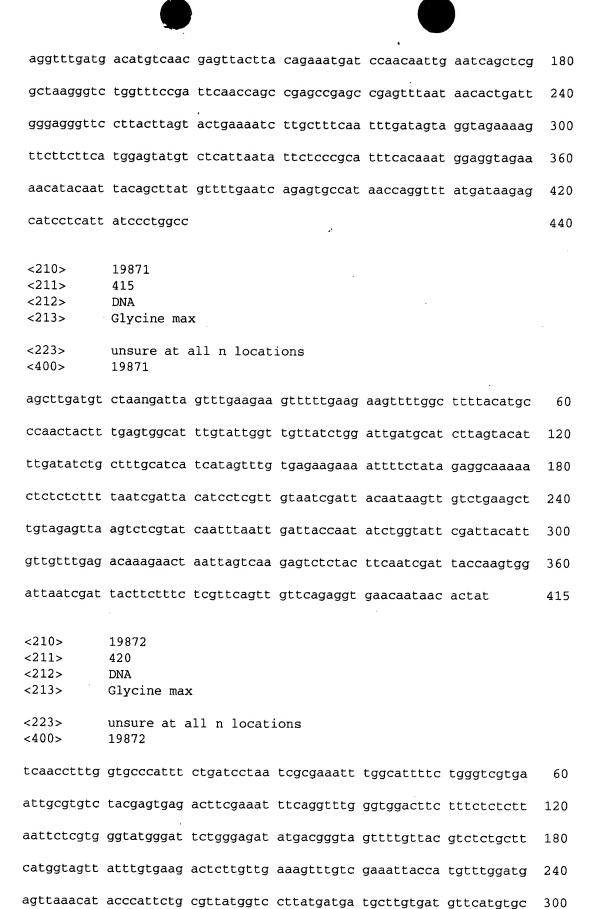


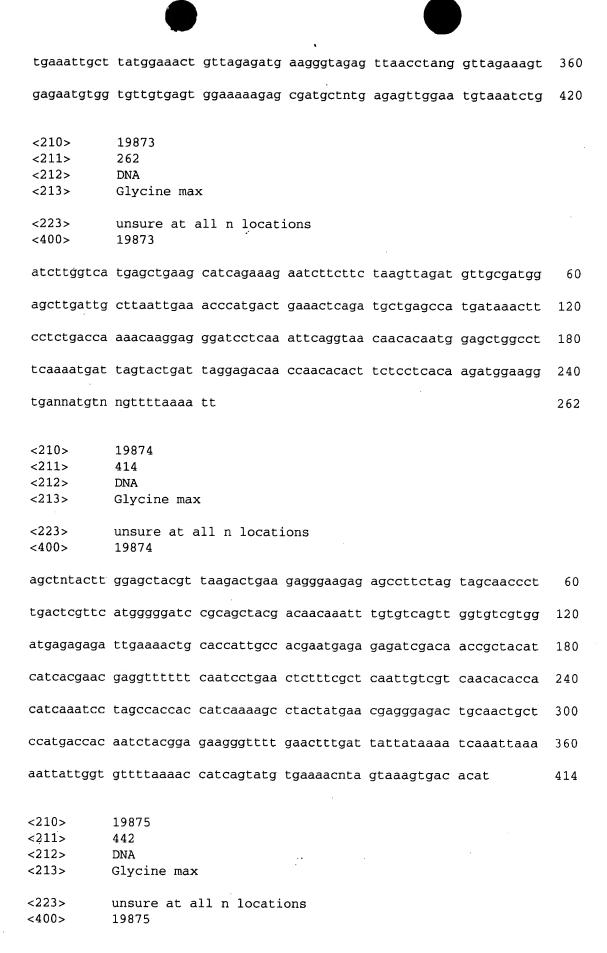
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|----------------------------------|--|-------|
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| acttatgca | c tnttatttta tatagaaact c | 391 |
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| cgaggcgctt | ccgtaacgtt teegtaaegt tteegtgagt aattaeacaa agattetega | 180 |
| ccgttcttca | a acattcatcg tttgttcttc gttttcttca gtcttcaacg ggtaagtacc | 240 |
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| atttttattc | togttttcat ttgcttttta taccccattt tgacgtgctt aagccattta | 360 |
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| atcttgacgt | gggagtgttt ctaaagagag aatgatgtaa gtatatttgg gagtgtattt | 180 |
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| tgccctaatt | gaatacacat tttgtcccga agcccttccc aagcaatgag tgtgtgttta | 300 |
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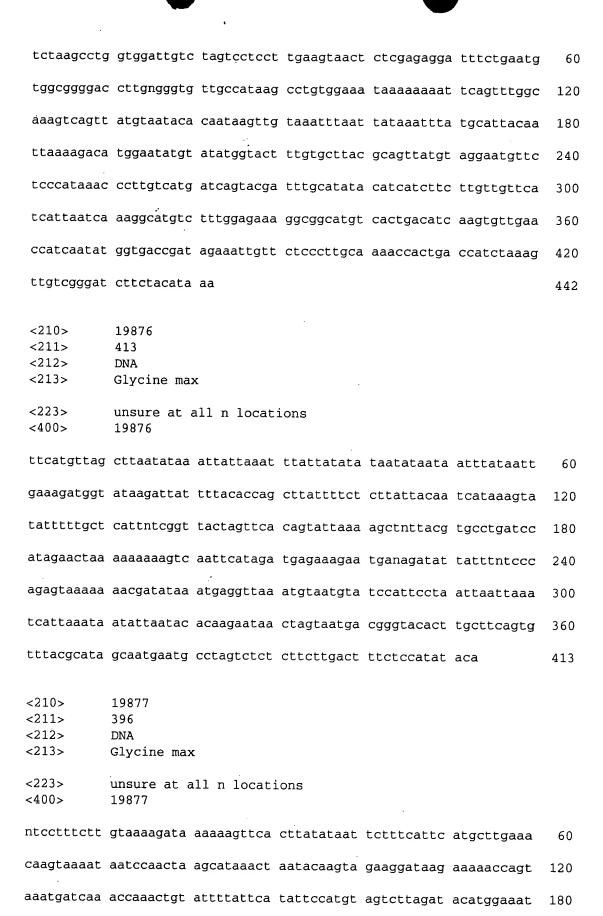
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| acctttttt | ttacttgtat acatgtaaaa | ttttaaatca | catcacttca | aatatgttaa | 180 |
| attagctcat | taaaaatata atttatttgt | aagtaaagtc | cagattgatt | atttactctt | 240 |
| ttatccaaat | tttaatttga aattcagttt | taaatgaaaa | aaaaggttat | tgtcactgac | 300 |
| tcattttata | aaataattct catccaaatt | taaattttta | atttgtaaac | tttaatcttt | 360 |
| nttttatcag | cctttanact ttaatctgaa | tatatactat | agtattttct | tttcagcaag | 420 |
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| cttcgtgaaa | gccaaatcac gatctctccc | atttgggatg | ggctttaaga | ttaattttgg | 120 |
| gcgtcccatg | ttctaaagca tatgcacata | tatagaaatt | gatcaataaa | gcaaaacttg | 180 |
| gggtcatacg | ttggggatag agagatcaaa | tagacaacta | gttcatgaat | tgatcgaatg | 240 |
| tttcggataa | gatgttgttc tagaaattta | aaataaaaaa | taattcaaca | aaaaagaaaa | 300 |
| tcatcaagga | tcaaagaaga atggttcaag | tctaatctct | cctctcatca | aggattaaag | 360 |
| aattntatag | aagaaaaagt aaactgtttt | tgaataatag | caaaaatgaa | agataaaagg | 420 |
| gaag | | | | | 424 |
| <210> | 19866 | | | | |

| | | 1.3 | | | | |
|-------------------------|------------------------------------|--------------|------------|------------|-----|--|
| <211> <212> <213> | 458 DNA Glycine max | . * | | | | |
| (225) | orline mari | | • | | | |
| <223> <400> | unsure at all n locations 19866 | | | | | |
| ntgaggattt | ggtctttgcc agtgaaagg | a tcgatgtggt | tctgaaaaaa | tgttaattta | 60 | |
| gtcatcctgc | ttggacgaat gagaaaact | a gggcaaatga | agagggtgag | aaagagggag | 120 | |
| aaacccatgc | tgtgactgcc attcctata | c gaccaagttt | cccaccaacc | caacaatgtc | 180 | |
| attactcagc | caataacaaa cctcctcct | acccaccgcc | cagttatcca | caaaggccat | 240 | |
| ccctaaatca | accacaaagc ctgtctacc | g cacttccaat | gacgaagacc | acctttagca | 300 | |
| caaaccaaaa | aacaccaaca aaaaggaat | ttgcagcaaa | tagcctgtag | ggttcacccc | 360 | |
| anattccgtt | gtcatatgct aaacatgat | c ccatatccac | tcaataattc | aatggtagcc | 420 | |
| ataaccccaa | ccaaggttcc tcaacctcc | a tttttctg | | | 458 | |
| | | | | | | |
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| <211> | 425 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |
| <223> <400> | unsure at all n location 19867 | cions | | | | |
| agcttccatc | aagttttaat caaagcacaa | a gagetteaag | taggtgctcc | ctaaacctcc | 60 | |
| attaagtttt | tgctttacct tctcttcta | tgttgtttct | tcatttttct | ccatgtatct | 120 | |
| cctcacatgt | cttgtgctaa atgtttttaa | a catgattctt | tagagtttcc | accgattaaa | 180 | |
| cttcctatag | aagctagatt tgattttcta | a tggttcaaat | ttcttgttct | tgttcttgaa | 240 | |
| ccataaattg | tgttgagttt aggttccttt | gagttttgtc | ttgttatttt | ttgtggctga | 300 | |
| aacctaaacc | ataaaattct tacaaaaata | a ttaaagtata | agaaaacctt | anaaatctag | 360 | |
| agtgacttgt | tcacctattg tagttntgto | c atagaagtca | tgtctagtca | tgaaacttgt | 420 | |
| cacat | | | | | 425 | |
| | | | | | | |
| <210> | 19868 | | | | | |
| <211> | 452 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |

| <223> <400> | unsure at all n locations 19868 | |
|-------------------------|--|-----|
| tatatgatat | aaactettat eettatgett teacaagtta ttttteaaag getattggaa | 60 |
| tttattttat | catctattat tcaatgtcca acacattatt attaaacaaa catgaaattg | 120 |
| tttggaaaaa | a ccagtactta tgtagtgagt gtctttacac gtacaattaa aaaagacata | 180 |
| actaagttac | actaatgttt tatattctgt ttgttaacat ccaaatcatt ttgcttgcat | 240 |
| gtgaaagcat | ctccaagata aatatttgct gatgcaatcc tccctaggaa gggaccagtc | 300 |
| actagagcca | tgagcaagag gctccaagag gattangcta gagttgctga agaatgccct | 360 |
| aagattetea | tgaacccagg gtagatntct gagcccatgg gccaaggttg agtccaatta | 420 |
| tctttgtaca | tattagacta gaatgtcatt at | 452 |
| <210> <211> <212> <213> | 19869 417 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19869 | |
| agtcttgttt | atattcaccc caattccaat gtcttatgct gacttgctcc catatctact | 60 |
| tgataattca | atggtagcca taaccccaac caaggttcat caacctccgt ttctccgaga | 120 |
| atacgactcg | aacgcaacgt gtgcttgtca tggagaagcg tcggggcgtt ccattgagca | 180 |
| ttgtaaggcc | ctgaagcgta aggtgcaagg tctaattgat gcgggctggc tgaaatttga | 240 |
| ggagaatcac | gtgtaaatcc tgacattgac aagagatgcc acacatggnn gaaatttgaa | 300 |
| agctgttgtt | agatgtctcc aatgactcat cangattttc aagtntgtac cattattgta | 360 |
| aaccacagtt | acaatggtaa atgaaatgga tatctttgtc cctcatcctc tcacaaa | 417 |
| <210> <211> <212> <213> | 19870 440 DNA Glycine max | |
| <400> | 19870 | |
| | gagaacatga agaactttct ttgcagacat ctatagggag cgatgtgaac | 60 |
| acacaggtac | ctatgggaat tttgcggtct gctccgagtc aactacacta gtttggcact | 120 |







| ttgatcacat | gaaçcccatt | attctagtcc | ctttntttga | cacaagtaaa | ttaacaagaa | 240 | |
|---|---|------------|------------|------------|------------|-----|--|
| acaacattga | gtgacactac | ttaattataa | caaacaaatt | aacggaataa | gtgatgacta | 300 | |
| gtactactta | ttagttgtag | tatgttcctg | gggtttggaa | ccagtgcatg | tatatttaat | 360 | |
| ttcctttctc | ataaatctgg | actttggtgc | ataagt | | | 396 | |
| <210> <211> <212> <213> | 19878 293 DNA Glycine max | | | | | | |
| <223> <400> | unsure at all n locations 19878 | | | | | | |
| ttgcttctat | ataagctaaa | ccattttatc | aagaaacaca | agttgagttt | tattcagaaa | 60 | |
| attagaggtt | atctctttta | tcttagtgag | agtgattctc | ctaagctctt | gagtgattca | 120 | |
| agaacatcct | gactatatca | aaggactttc | acaacctttg | tgtgttgccc | tcgccggaaa | 180 | |
| gagtgattct | ttccttcctt | tcaccttcaa | ccttttttct | ttcaaaccac | aattccagaa | 240 | |
| aatccacttc | tgcccagaat | tatcttgtgg | ccatganctc | tcgtttacgt | gct | 293 | |
| <210> <211> <212> <213> <400> | 19879 130 DNA Glycine max 19879 | : | | | | | |
| gtgtcaataa | gtagttgtag | tttctcatta | tggggaatac | ttgtacttgg | gggcatgtca | 60 | |
| ctcggttcag | aactttcttg | agactcatgc | tgatcaccat | aaaaaaaaaa | gagttgtctg | 120 | |
| ggcccgacat | | | | | | 130 | |
| <210> <211> <212> <213> | 19880 400 DNA Glycine max | | | | | | |
| <223> <400> | unsure at all n locations 19880 | | | | | | |
| tcttgctntg | tcccttgctc | gtgcctttag | ccatcatctt | actttgatgg | gtatagctga | 60 | |

aacccaccat aggtatacat tcttgtctcc aactattgtt gtctttttat tccattagtt 120

| gataatttg | t cttgtcttta | tttggctgag | ctctgtgcaa | a aattaagcc | g caatctctcc | 180 |
|-------------------------|------------------------------------|--------------|------------|--------------|--------------|-----|
| aaaataagc | c gaacctttga | ctaataacaa | cattgaggga | a atactatacı | tctttaatct | 240 |
| aagattacc | c cagtatgata | gtcttgaaca | agttatgttt | ttgcagggtt | cgtaaaggag | 300 |
| gacatatgg | c tcttgctgca | aaatcttgtg | atgatatctr | ı taacnaacct | gttttataag | 360 |
| acagtgttc | a agcacgtctg | tggtttagtc | ctcatttctc | ı | | 400 |
| <210> <211> <212> <213> | 19881 431 DNA Glycine max | | ions | | | |
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| tggtgattag | gatgttntga | ctgacttcta | ctcatagatt | cattccagtg | ctcatggtga | 60 |
| tcctcctcat | cattaccttc | tgaagagaaa | gaagtgatca | tcaagtctta | aaagtggatt | 120 |
| gaaagaatat | caaagatatc | tttgaggata | taacagtgac | aaggcgtagt | ctagaagaac | 180 |
| aaatcatata | gaaaaatctt | cagctaatca | tgtaaagaca | ttgtgagtgc | attaatagaa | 240 |
| caataaatgg | atgaccattt | cacttgcagt | gtatcatcaa | atactatagt | gtatctatat | 300 |
| tagtctaact | cacaccaagg | tttaaaatac | tttgtagtat | ctacaaacta | ttttcatgcc | 360 |
| ggaagaaaac | tttncatttg a | aaaagcagat | tntgttgcct | catcatatga | aaccttntgc | 420 |
| aaaaacattc | С | | | | | 431 |
| <210> <211> <212> <213> | 19882 499 DNA Glycine max | | | | | |
| <223> <400> | unsure at al 19882 | ll n locati | ons | | | |
| cgccacaccc | actcaaaacg a | igccaaacac | aaaacgttca | aactaaaana | aacccccncc | 60 |
| cgcccaagga | tgacacantg a | laaccctgaa : | naccaccaaa | tccagcgccc | ccacaagacc | 120 |
| caacaagcca | ccccacaata t | tacacccat | ttacacccac | cacaacgggg | aaacggggac | 180 |
| gaaacgagca | cccacaacca a | .cactaacac 🤉 | gcaaaaccaa | accaagaagc | accacccaca | 240 |
| acaagaccaa | aaaccaaccc c | acacacaac a | aaaaccgaa | acgaccaccc | ggccaaccca | 300 |

| cacaaaaaa | c caaccccaa | a cgaaacgcg | a accacacac | ca caacgaacc | c aaccacaca | 360 |
|-------------------------|-----------------------------------|--------------|-------------|--------------|--------------|-----|
| accaccacg | c cagcgacaa | c acagacacc | a accgcaacc | c cacacaaca | c aacaccacaa | 420 |
| cgaacccca | a accaacaca | a caccacccc | c aaccaacco | a gcaagcacg | c accccagaca | 480 |
| acaccaaca | c caacaccac | | | | | 499 |
| <210> <211> <212> <213> | 19883 406 DNA Glycine m | ax | | | | |
| | | | | | | |
| gtgcgtagco | caccatctt | t ccataaaagt | atctatacg | c gtcttccate | c acgatcatcg | 60 |
| actccctato | : catcattgg | g ggtaccacct | gggccgcta | g atccctccad | cttttgggcg | 120 |
| tgttctttgt | aagatctgto | tcactttttg | g caaatgttc | t gtagttgcat | cctatcctga | 180 |
| accatatcaa | acttgttctg | g atactgccta | tcaaaggcca | a ccattatgto | cttccaagaa | 240 |
| tggactcggc | aagacttcca | gtcagtgtac | caggtaacag | g ctaccctagt | aagactttct | 300 |
| gggaatgaat | gtatcagcaa | gtcctctagt | tctcgtatat | ccccatctcc | : tgacatacat | 360 |
| ctttgatggt | cttgggaaag | agtccacttt | acttgtcata | ı tcaatc | | 406 |
| | 19884 424 DNA Glycine ma | | | | | |
| <223> <400> | unsure at 19884 | all n locat | ions | | | |
| agcttgagaa | ggaacagaat | tatttttgta | tccactctca | gctctgtagt | ttaaaccaaa | 60 |
| atccattcc | aaacaagcac | aaagaataaa | aaatcattgt | aattattagg | tctgttgttt | 120 |
| tgaggatat | cctctcccta | gttggattgt | ccttttctat | ctaacaagct | tactatttta | 180 |
| ctgaaaata | aaagcacttc | ttggtatttt | actatttagt | atttactata | ttatacaacg | 240 |
| aaagaaaag | catcttaaaa | ttattcataa | aacttactac | aataaacatc | atcatcattc | 300 |
| acanaaaca | ttcaaaaatg | aacgagtcaa | caatgtgaag | tttgggaggc | caaaatgaaa | 360 |
| atgaaaaat | gcaattacag | caaacatgct | taaatccaaa | agaatctaga | acaatactga | 420 |

| catc | | 424 |
|----------------------------------|--|-----|
| <210> <211> <212> <213> | 19885 451 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19885 | |
| ntgaatcata | a aaaaaaatac tttaaatagt aaaattttta tttaaagtgt ttttcaacaa | 60 |
| aaagtaaaa | a gaatatattt ttataataat tttttataaa aacaaaataa aataataaac | 120 |
| actaaattad | c agtaagtaat atatatata atatatata atatatat | 180 |
| atactccttc | c caaagttgcc actggtaatc tgattcctgt tccatcttct caatatccac | 240 |
| tcaccctttt | gtgctgccac taacaaaaca gcacttcaac ctcaaatttt caacctttgc | 300 |
| tcattcatgt | controlled the control of the contro | 360 |
| tagtcattct | cgcaaccagg ttagtttcat gtctttcact tcccatatct ctcttcttt | 420 |
| gtgtttttg | ttccaacaac taagcaaaat a | 451 |
| <210> <211> <212> <213> | 19886 399 DNA Glycine max | |
| <400> | 19886 | |
| agcttggttt | ttcttattcg atattttaa tttgcaaaat tgaatgttat tggagttgca | 60 |
| ctgtttatca | tgcttgtctt gtacttaaaa atatttaact tctgcatttt ttttattcca | 120 |
| cagaactttt | acatgatagg aagggacaag agcaggacat attggaaagt actaaagatt | 180 |
| gaccgtcttg | atccttccga gctaaatttg cgtgaagatt ccaccacata tacagaaagt | 240 |
| gaatgttctg | atcttttgag acggatacat gagggtaaca agtccacagg tggactaaaa | 300 |
| tctgctacaa | cttgttatgg aattgtaggt atgtaaatcc ataatgtctt agctacctgc | 360 |
| ctgttgacca | attatgagtt actgtctgga tatactatt | 399 |
| <210> <211> <212> <213> | 19887 301 DNA Glycine max | |

| <400> | 19887 | | | | | |
|-------------------------|------------------------------------|-------------|------------|------------|------------|-----|
| tggatgggcc | ctgaatattc | tagctctaca | gcttgtgcta | caagtaactt | ttggatattg | 60 |
| aatgcacgca | accatatcat | tagaggtgaa | gctgcaagta | tttttgcttt | gatatgtgct | 120 |
| cttaggatct | taaaactctc | atattaaccc | tgacctttac | ttattgggac | aaaatttggc | 180 |
| tcaatgatgc | tctctctttg | tcaaaaggcg | tcaacaacca | agtaacatga | aatctgctgc | 240 |
| cgatggaagc | tctgggtggc | cagagctgat | atataccatt | ggatccttta | tactattctc | 300 |
| t | | | | | | 301 |
| <210> <211> <212> <213> | 19888 417 DNA Glycine max | | | | | |
| | | | | | | |
| agctttgctc | gtatttgtca | agtgtatgga | ccacgttgta | gccaaggtgc | tcatcgataa | 60 |
| tggttccagt | ttaaacgtga | tgcccaagag | cactttggag | aaattacctt | tcaatgcttc | 120 |
| ccacctaaag | ccaagttcca | tggtgtttcg | tgccttcaac | gacacccgcc | gagaggttag | 180 |
| gggagagatt | gacctccccg | tacagatagg | ccctcacacc | tgtcaagtta | ccttccaaat | 240 |
| aatgggcatt | aaccccctt | acagctgcct | gttggggcgc | ccgtggatcc | actcggtggg | 300 |
| agttgttccc | tctacactcc | accaaaagtt | gaaattcgta | gtggaagggc | atctggtcat | 360 |
| cgtatcaggc | gaggaagaca | tcttggtaag | ctgcccatcc | tctatgcctt | atgtgga | 417 |
| <210> <211> <212> <213> | 19889 444 DNA Glycine max | : | | | | |
| <223> <400> | unsure at a 19889 | ll n locati | ons | | | |
| ctaagcttga | caaaattacc | actgctgtct | tctaacaaga | tttaaaggga | aatctctcta | 60 |
| aagatacatc | taataatacc | cataattata | acacatacaa | aacactgaca | gaaacaatgg | 120 |
| gtatccaatt | caagaaggat | acaactatta | acaaaacaaa | tattcacaaa | taataaaaga | 180 |
| ataacaaata | gacacttaaa | gaactaaaat | aaacttccca | aatagaaaga | agttcctcgg | 240 |

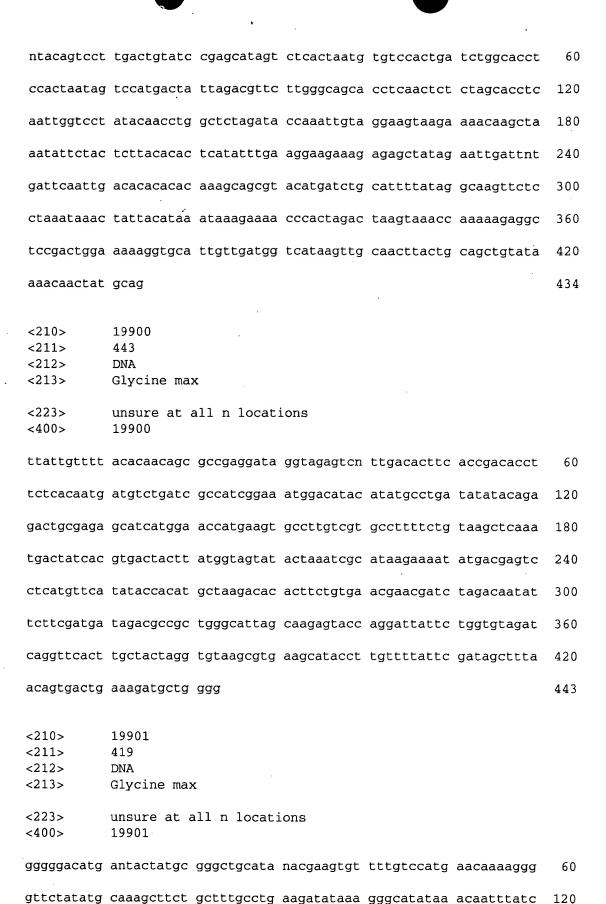
| caacggcgcc | ataaacttgt | tcgacgaccg | acaagtgcac | : cggatcacgc | aagtagtata | 300 |
|-------------------------|------------------------------------|--------------|------------|--------------|------------|-----|
| acacggtaag | tgaataccga | gtatcgaact | ctcgaggaac | : ttgttttact | tggtaaagct | 360 |
| gtggttagta | aataagtgtt | ttgggtgaaa | cttgtgtctg | gtätgacaag | atgcanacta | 420 |
| actatcaaaa | gaaatacgtg | agta | | | | 444 |
| <210> <211> <212> <213> | 19890 409 DNA Glycine max | ζ | | | | , |
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| agcttatgtt | atggatcact | gtatggtttc | ctttaacctc | agtctatcca | taacgcaaag | 60 |
| aacaaacatg | aagaatgcca | acattacatt | caaaaatggg | tcaaggaatc | acaacgaana | 120 |
| gtgtacttgg | gaccttacct | aaatcagtaa | gttgaattga | tgttgtagaa | tatagatatt | 180 |
| atgtgcatta | ttgttgccta | actaatgttt | ttcgtcttca | aggcacattg | gcaacttgtt | 240 |
| gttctgtgtc | cacgggacaa | tattgttgtt | tggttttgtt | ctttgtgtaa | gaagcctgat | 300 |
| gttaacatca | agacaacaat | gaataagtta | gttntaacat | taaaagtcaa | ttgaaatatt | 360 |
| gcaattgtan | ggtataaaga | caatgattat | ttgaatatat | acgttaacg | | 409 |
| <210> <211> <212> <213> | 19891 350 DNA Glycine max | | | | | |
| <223> <400> | unsure at a | ll n locati | ons | | | |
| ctcagcttct | cccccaattt | ctataatagg | gggagaagtg | aagtttatna | gggttcagcc | 60 |
| ctcctggtaa | ttaaaatcac t | ttanaattag | tgagaaaaat | tggtttcgtg | aagaaaatcc | 120 |
| aagccgaggc | gctttcgtaa (| cgtttccgtg | gatgatttcg | cgaaggtttt | cgaccgttcc | 180 |
| tcgacgttct | tcattcgttc t | tcgccgttc | ttcggtcttc | aatccggtag | ttccctagat | 240 |
| cgaacttttc | aattcattct a | atgcaccctt . | agtggtcctc | atttgttttg | cgtgctttca | 300 |
| tttacatatc | atttactttt d | eggeceett | ttgcatgctt | aagtcatttt | | 350 |

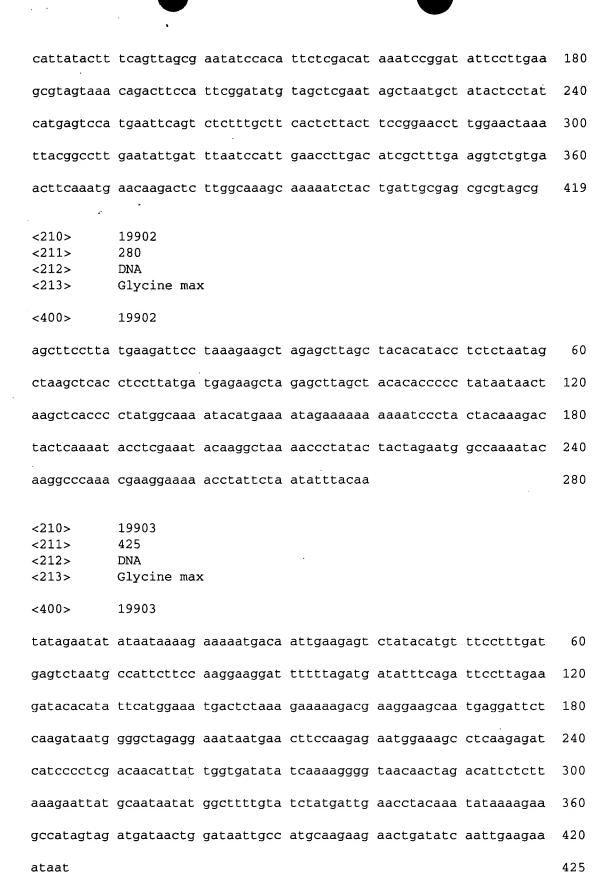
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|---------------------------|------------------------------------|--------------|------------|------------|------------|-----|
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| tttgtctgca | gactataata | ggaagcaacc | agattatata | aataagaaga | attatggcaa | 120 |
| aaagataaac | gagtaggaat | taacacgagg | cgcattgatt | ttgaacccgt | agcaaactga | 180 |
| atgcaccaga | cacttatgca | accactggaa | attgcactac | acatgatata | taaacacatt | 240 |
| aataagctaa | tcagcatata | attatggaga | aaattcattc | tttacttgg | tatggcacct | 300 |
| tgtcgggaaa | tatg | | | ٠ | · · | 314 |
| <210> <211> <212> <213> | 19893 363 DNA Glycine max | ĸ | • | | | |
| <400> | 19893 | | | | | |
| tcgttgtgat | gctagagctt | agctactttc | tcccatgttt | taactaagct | cacctgcttg | 60 |
| agaagctaga | gcttagctac | acacacccct | ctaataacta | agctcacctc | cttaagaaga | 120 |
| gaagctagag | cttagctaca | cacccctata | atagctaagc | tcacccccat | gactaaatac | 180 |
| atgaaaatac | caaggaaaaa | tgctactgca | aagactactc | aaaatgctct | gaaatacaag | 240 |
| gctaaaaccc | tatactgcta | gaatggccaa | aatacaaggc | ccacaagaag | aaataaagcc | 300 |
| tattctaata | tttacgaaga | agagtggagc | caaccttgac | ccatgggctc | agaaatctac | 360 |
| cct | | | | | | 363 |
| <210><211><211><212><213> | 19894 417 DNA Glycine ma: | × | · | | | |
| <223> <400> | unsure at a | all n locat: | ions | | | |
| agcttgattt | aagaaaatta | ttaattaatg | taaggttaca | aaataagtag | ggataattaa | 60 |
| ggttgattaa | tgacgatcta | gatttcatgg | aattagaaaa | tgggtaatta | agtcacaaga | 120 |
| gtttaaaatg | gagggcattt | ttgtaaatga | ctatacaact | agtttaaaaa | tagaaattta | 180 |

| gtttaattag | ttggtgacaa | a attaaagtgo | ctgattatac | : aatgtagaat | aattaaaata | 240 |
|-------------------------------------|---|-------------------|------------|--------------|--------------|-----|
| | | | | | actctgtgtt | |
| | | | | | | |
| gtgtcatctg | r tgcatgtant | : gaattaattt | aagtatttat | atgctnttaa | a tcatagaatn | 360 |
| ttgtgttatg | tatatatgtg | ı tgtgtgtgtg | tatgtgtgtg | gttagttggt | ttaataa | 417 |
| <210> <211> <212> <213> <223> <400> | 19895 433 DNA Glycine ma unsure at 19895 | x all n locat | ions | ¢. | | |
| tatcacccgt | atcagcgatt | gaagaaggtt | ttaatgggtt | tcaggaaaat | gagagtgcgt | 60 |
| | | - | | | actatctttg | 120 |
| | | | | | tcaatattct | 180 |
| | | • | | | atgcatgtgg | 240 |
| | | | | | cactcttctt | |
| | | | | | | 300 |
| | | | | | accacaaaaa | 360 |
| aaaaactatg | tttgctgata | aaaaaataga | aatcaaacag | caattggttt | gctttattnt | 420 |
| ttcccatcaa | tca | | | | | 433 |
| <210> <211> <212> <213> <223> | | x all n locati | .ons | | | |
| <400> | 19896 | | | | | |
| agcttaatat | tcactaccat | ataatatgtc | tggacatata | attatatgat | aaaacttgtc | 60 |
| tttgagctta | ataagtactt | tgcgctcaaa | taatcttcgt | tgttttttt | catttaagtc | 120 |
| acccgactat | atcctatgtg | tgacattctc | attaatttct | tcgtcatctt | gtataattga | 180 |
| tcctagatat | ctaaacttag | agacatgtga | tgtgagatct | tcccccaatt | ttattaaagt | 240 |
| ttcaattaca | tttttttct | aatttaagga | ctttaatgaa | ttacactttc | ttatgcagga | 300 |
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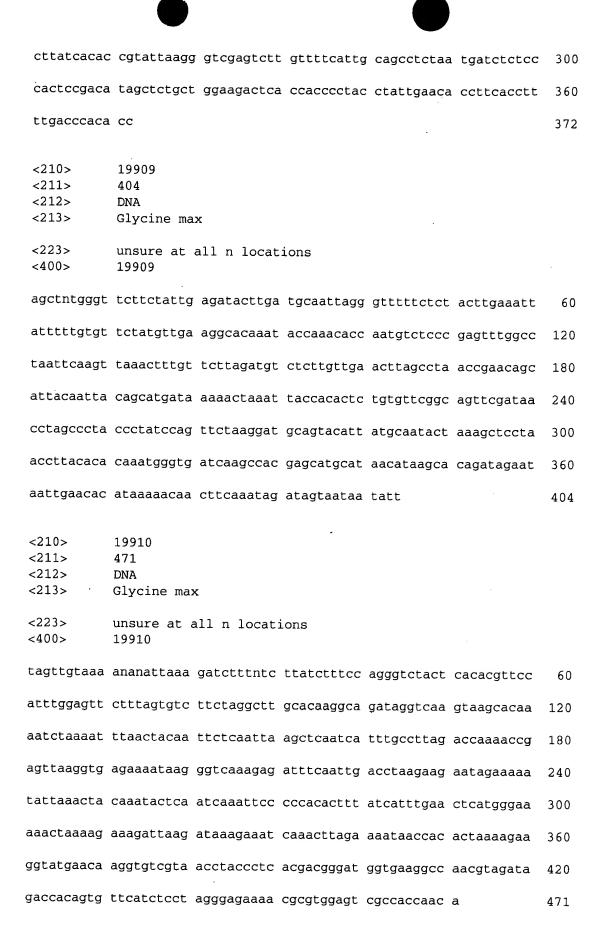
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| | 19904 400 DNA Glycine max | | | | | |
|--|--|--|--|--|--|--|
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| tactggtaat | cgattaccaa | aacattgtaa | tcgattacag | ccttttgaaa | ataattggaa | 120 |
| cgttgtaaat | tcaatttgaa | aactttntca | aaacaatntt | gctactggta | atcgattaca | 180 |
| acaatctggt | aatcgattac | cagagagtaa | aaactctttg | gtaaaaggtt | ntgtcaaaaa | 240 |
| ctcatgtgct | attcaaagtt | gtgaacaact | ttntaatact | tatcttgatt | gagtcttctc | 300 |
| ttcatccttg | attcttgaga | tcttgaacct | tgaatcttga | ttcttggctc | tagactttct | 360 |
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| <400> tgtaatgagc aattcaaggt caaatatgca aatcattggg aacttatcac aattactcat | 19905 ttcataacaa aaatccattt tgtgaattaa gcaaaacaca ttccaaattg | agaaagggaa ggctagaggc gaagcatcag ccaaatgatt agctttcaaa tatcctataa | acataggctt ttataagaac caagaatcaa atgatgatgg actatcataa ttcaaagaan | aaaattgcct gccaaggcta ctcaaattct gcgactttta aatatngcaa | aaatcatttc ttgtgcaagc cacaaaggta ttttcaaaac agtgtacaag | 120 180 240 300 |
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| atctctcaca | tacggataaa | agaagttgaa | ttcctcatcg | aaagtctctg | tagtataatt | 120 |
| cattatctgc | atcactccta | agtcaactca | nacttattta | tgaaataaat | caagttaaaa | 180 |
| ccttntaaaa | atctatttag | ttacaaaaac | aaaaacagtt | ccaaactata | ttaaaaaaag | 240 |
| ccgaaatgat | cgacgaaggt | ttgttaaaac | tagaaaatta | gtaaaataaa | tcttcgaaca | 300 |
| tatgtgacaa | gtataggttc | tgtcaaaaag | aatagtataa | atacagta | | 348 |
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| tgagaatagt | acattatagc | attaaatctt | gattattatc | ttatgcagat | gataaattgg | 6,0 |
| ttttaaataa | gataatcccc | gtacacacac | atgataaaag | aatagaaaac | attgtgctnt | 120 |
| gtagaagtac | taaaaccaca | ggaaaatata | gagaagctag | gactaagaga | atgaaagcca | 180 |
| aagtaccaaa | gttttgtatt | ataaccctgt | atttctttac | ccttcattta | ctttttgcca | 240 |
| ttttagtaga | acttttgaga | tatgaggatt | atttttaaca | ctacacttta | cagcaacagg | 300 |
| tatttcgcag | tacaaagtgt | tgtcctgttc | tataaggatt | actgaacaca | ggaaagcatg | 360 |
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| agctntacaa | cttgcttctc | tacctcacgc | cgtcactctt | gcctgactgc | aagaggacaa | 60 |
| gattaatgac | cgccgaaaac | actctcggac | ctcgtttcct | tccccacaac | aaccagcttc | 120 |
| agcaccatcc | actcaaaccc | aaacacctcg | ccctaagact | cctttcatac | agcgtactca | 180 |
| cgaagagatg | acctatatgc | gcgaaaaggg | cctctgctac | aactatgatg | agaaatggaa | 240 |



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| taagaagata | ggcctcacag | tgatcaagaa | tgaaaaggat | gagcttatcc | ccacaagagt | 120 |
| gcagaacagt | tggcgagtct | gcattgatta | taggaggctg | aattaggtaa | ccagaaaaga | 180 |
| tcattttccc | ttgcctttca | ttgatcaaat | gcttgagcgc | ttggcaggta | agtctcatta | 240 |
| ctgctttctt | gatggtttt | ctggttattt | acaaattcat | attgctcttg | aggatctaga | 300 |
| aaagaccaca | ttcacctgtc | cctttggcac | ttttgcctat | atgaggatgc | cctttagcct | 360 |
| atgcaatgcc | cctggtacct | tccagcggtg | tatgcttagc | attttcagtg | actttttaga | 420 |
| gagtcgcata | gaggtggtta | tggatg | | | | 446 |
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| atgtttatat | gctgaaattg | cctatggaaa | ctgttagaga | tgaagggtag | agttaaccta | 120 |
| gggttagaaa | gtgagaatgt | ggtgttgtga | gtggaaaaag | agtgaggctt | tgagagttgg | 180 |
| aaggctaagt | ctgaattctg | tggtaaatgg | aggttaaaat | gagttaatcc | tagcttgaaa | 240 |
| tgtcatttac | aacatgtgac | aaaggttagg | ttgtgctata | gggaaaaact | aatgaccaaa | 300 |
| gtgaacaaag | agccatttct | ggggcacaat | tgggtgttga | atagtcaaat | tttgattcgg | 360 |
| tggaatttta | ggtgtaaata | cagtatgggc | aagtctatat | tgatgttatg | gactggtgtg | 420 |
| aggtgagagt | ttgcctcata | tctacctcat | tctaaatctc | acc . | | 463 |
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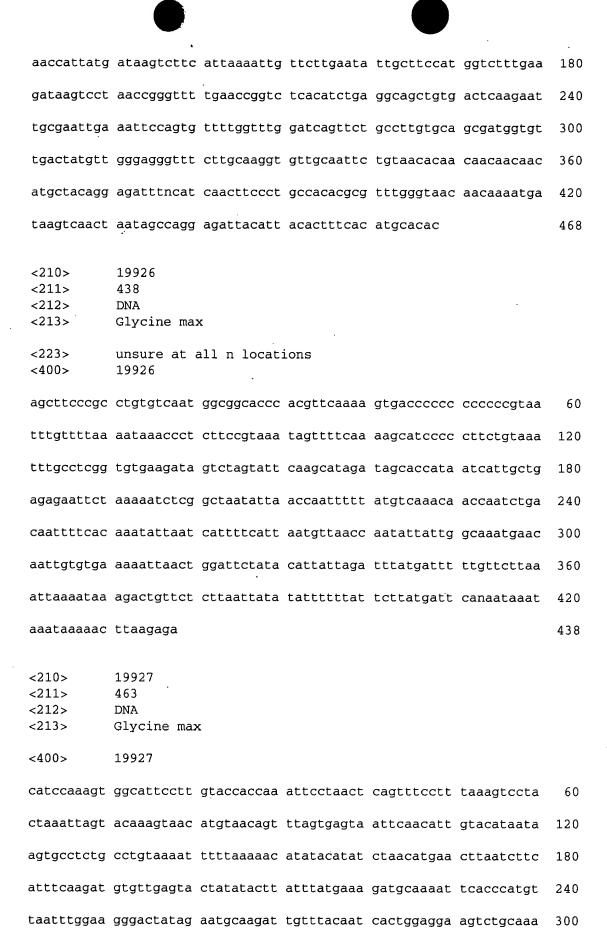
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| tttcccttt | c cttgttttga agctcactac aagccttaag tgaaaaacca tgatatcacc | 120 |
| atatcctta | a ggaattttgg agctttggaa ttgttttggg aataagtgtg gggggttttt | 180 |
| gtttcattg | g ataacttgtt ttgttggcta tgcttcatga tgtattttgg gccatacttg | 240 |
| atgtacacto | g catattggtt aaatgttgga catgctgaat caaatgttgt ttctcaaagg | 300 |
| ctatagagta | a aaaaaaaaaa aaaattcaaa aaaaaaagag aaaaagaaaa gcaataaagt | 360 |
| tgagtgaata | a agatettana tggcacaaga atcatgaaac tetttggtat actetetatg | 420 |
| tctaaattnt | atctttactt ctt | 443 |
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| ggcacttctc | tetetttega atntgettgg aaaaattget teegtgaaga aaateeaage | 120 |
| cgaggcactt | ccgaaacatt tccgtgagga atttcgcgaa ggtttcgacc gttcttcgac | 180 |
| gttcttcctt | cgttcttcat cgttcttcga tcttcaacgg gtaagtacct cgaaccaagc | 240 |
| ttttcgattc | attctatgta cccgtggtgg tccacattgt gtttcgtgta tttttattct | 300 |
| cgtttcattt | gctttntata cccccttttg acgtgcttaa gccattttat ttaagtcatt | 360 |
| tctcgcttaa | cctaaaaata aaataaattt ccaccgatcg tttgaattta ttatccgtta | 420 |
| acttcngtta | aatg | 434 |
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| gattntaaag | , atccttaatt | atgatagana | taaaaaaaat | ccttaattat | gatagattnt | 120 |
|----------------|--------------|--------------|------------|------------|------------|-----|
| aaagacaata | ı tgatattaca | ctnttagtgc | ataattattt | ttctcgaagt | aaattaataa | 180 |
| taataatctt | aaaaaatagg | acttattcaa | aaatgtttt | ttagttttta | gttatgaaaa | 240 |
| anagaccaaa | ttcaaattaa | aaaaaattga | aaaaattgaa | natataatta | aacttaaagt | 300 |
| ttataataac | acaacagtag | tatataactt | atantttccc | tctcgtctta | ccaatcttat | 360 |
| aaatatattt | tagaaatttt | tttatcatat | cgttaatccg | tgtcataaag | atccaatgac | 420 |
| tagtgacctt | t | | | | | 431 |
| | | | | | | |
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| <211> | 288 | | | | | |
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| \Z13 > | Glycine ma: | X | | | | |
| <223> | unsure at a | all n locati | ons | | | |
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| atttaaaatt | | . | | | | |
| gullyayall | ntctcactgg | taatcgatta | taggattcta | gtatttgatt | acatagttat | 60 |
| atgtttgaag | agttatgact | tttcaaaggt | tttttttaa | aatctcttca | atggttatca | 120 |
| attacaggat | tctagtaatc | gattactgat | cgaggccata | cccgaatcaa | ataaacatta | 180 |
| aaaatacagt | atctaggaag | tgatcctagg | tegteteeeg | acgagcaagg | gtcaaacaaa | 240 |
| cgttcataac | agatagtagg | aaaatattaa | cgaattgggg | aaaäaaaa | | 288 |
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| <211> | 195 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |
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| tgagcccatt | tagaggtaat | agataagatc | atcgcggcta | gaatggacat | gtgtagagat | 120 |
| actcacagga | tcagatcgcg | atttattctg | ggatatctat | tgtattgtga | tgcttcctta | 180 |
| atgatcatta | ataca | | | | | 195 |
| | | | | | | |
| <210> | 19918 | | | | | |
| <211> | 261 | | | | | |
| <212> | DNA | | | | | |

| <213> | Glycine ma | ax | | | | |
|-------------------------------------|---|------------------|-------------------|------------|------------|-----|
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| aactaagctg | gggatggctg | g cattcgacaa | agadggaagt | gcccactact | ttgcttgata | 60 |
| aaacaatggc | tagaccacaa | ı cagegetgga | ggcggcaacg | gacaagggac | tgacaaaaaa | 120 |
| attatgtaga | catgaacaaa | . caatagatca | tgcgcagagc | gtgccaggtg | aaccaagaga | 180 |
| agcactgaac | aggtgttaga | ccagctacaa | acaatgtgcc | tgagcaaaac | gaaaaccaaa | 240 |
| aacgtgaccg | acacagatga | t | | | | 261 |
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| | | aaaagtttaa | + 222+2+++ | | | |
| | | | • | | | 60 |
| | | aacattttaa | | | | 120 |
| gtttatttac | ggaacagtca | actaaaattt | ttaattatta | aaaaaatcta | gaagctaatt | 180 |
| aaaatatatt | ttgaaacata | acctaactaa | atcaacatat | ttttagaatc | ttgatttagt | 240 |
| ttgtatcata | tttgaatttc | tcttaaaatt | aattttatta | cttatgtttt | ggttaaattt | 300 |
| gagtctaaat | aagaagcgat | cataaatggg | ctgctcttaa | agtattcttt | atttatatan | 360 |
| ttacaataga | aacatgaaat | agaaataatg | ttaattggta | ttatctaaat | aattatcatc | 420 |
| ttatcaacat | ctttataaca | aa | | | | 442 |
| <210> <211> <212> <213> | 19920 352 DNA Glycine max | × | | | | |
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| cattacagca | tataatatat | tattacattt | ctgaaacgac | atacaacttg | tggtgttacg | 120 |
| atgggtgcat | agccaaggcg | atcctttgct | tttgagcaat | caaatgttct | gctgcaagat | 180 |
| atgagtctta | ttcttgaagg | aatgagttga | ggcagctcca | tcccatatgg | gcctagcaac | 240 |

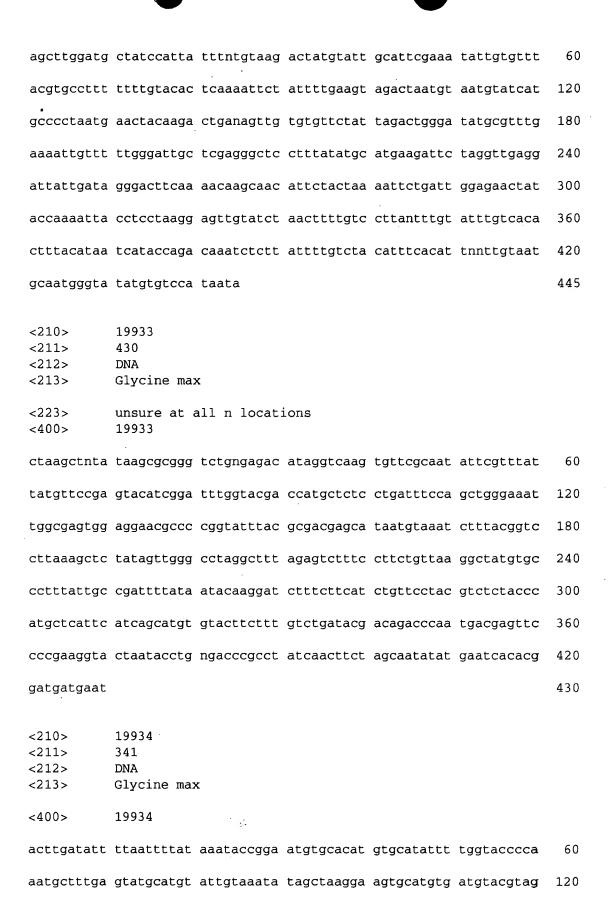
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| ccttggcct | a taagcaacat | gacttagatt | acaaagaaat | ggcattgaca | at | 352 |
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| ttgcatgcaa | gcttctactt | tagtgcttgg | cgggctgcct | tcactttgtt | gtctcgtacg | 60 |
| cgagctttga | ccactattct | tccttaccga | gatgcttctt | ttcatctcct | gctgagcggg | 120 |
| cttatagcct | ataccatact | tgccatgata | tacttgcggt | tttataacgc | tagttatgcc | 180 |
| gtcgttgtct | ctgtctaaac | ccatcccatg | ttcgtaaccg | ttccccaact | taactcgggc | 240 |
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| gctgaccacc | tcaa | | | | | 314 |
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| agcttagaat | tattgagata | tcatctgtgc | gtttcatgga | cccaggagcg | gactttgact | 60 |
| tcagtttctt | aattcttcaa | ttgtatgttt | aagtttgaaa | caatttttt | ataaaaaaaa | 120 |
| agagggaaaa | gacatcaagt a | aacgcaattc | tccttggaaa | aaagtcaaga | ttctgcacta | 180 |
| ggtctacata | attcatataa a | actattaagt | taaatgacaa | aatatataaa | attatgaaga | 240 |
| aacaccataa | tataataaaa a | agagtaccac | attcatacat | accctcatca | acatcaacag | 300 |
| caaccacaag | tttctgggat a | aatcgatgat | cactgaaaca | aggaggctta | gcaggacttc | 360 |
| catttataga | tccattatca a | accttaacca | gagtagcatc | actctcatca | ctcctaac | 418 |
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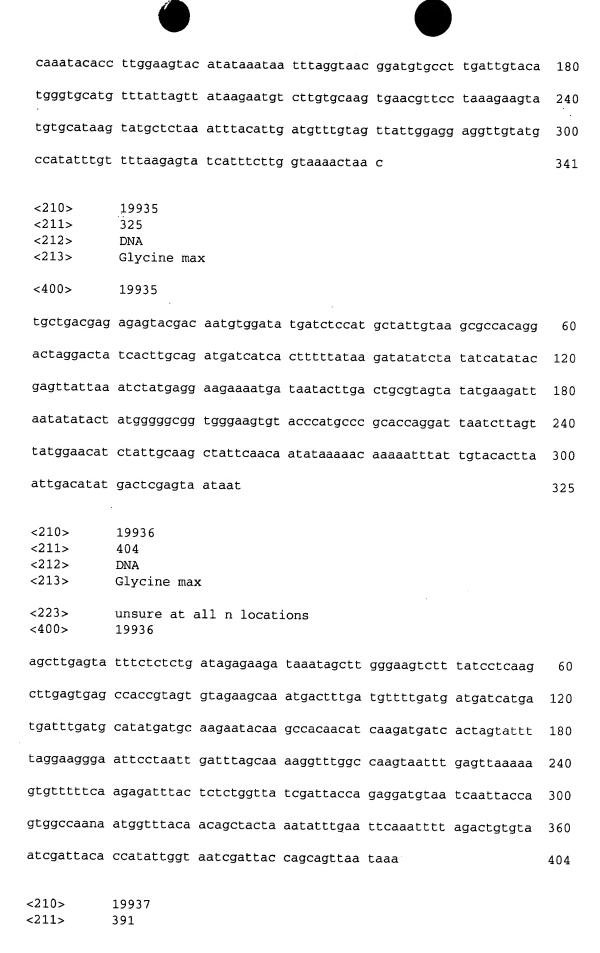
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| tcgattgaa | c tcgtatatgc | acaataatgt | ttaatgagaa | a gggaagagac | aganagcaaa | 120 |
| acgtaatca | ttggattaca | gtccatggtt | caaacttcag | g agttccagag | acaggaacct | 180 |
| gaatctacti | gcaaaaaata | attaaatgaa | aactgaataa | ı gctcagccaa | tgttgtacaa | 240 |
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| tctggcaato | tcttttcctt | tctcagacca | acgatgcatg | tgcgatatag | gagcagaaaa | 360 |
| aattattgaa | ı agaatgatga | tactagaacg | ttctatcact | tatataccan | atcatggcaa | 420 |
| ggtttgtcac | aa . | | | | | 432 |
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| ttgtgatatg | gtattgtaaa t | tgacatctct | ggctagagtg | gctttgatta | ggatgacact | 120 |
| aagtacatga | tcacagttga (| gaataaaaat | gcttggaatg | aataatgcat | tgtaagtatt | 180 |
| cttttatata | tttctatctg t | ttattcacaa | ctgattagat | ttgactttgt | ttccagtcac | 240 |
| ataaatcggc | taaaccgttt d | cgattcaaag | tgcttcacaa | ctgggatgat | atagtggatt | 300 |
| tgtgcgctaa | agatagagcc a | actggtcatg | gagctaaaac | tgctatggat | atttatgaag | 360 |
| cgatgagttg | agaaa | | | | | 375 |
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| ctataatact | cagcttgagc c | aaagggaaa | aagaacttag | ttattcaatg | tactctcctt | 60 |
| aatanagtgt | gttcaatgca a | agttaccac | tctattccat | ccaaccttga | ggattgataa | 120 |



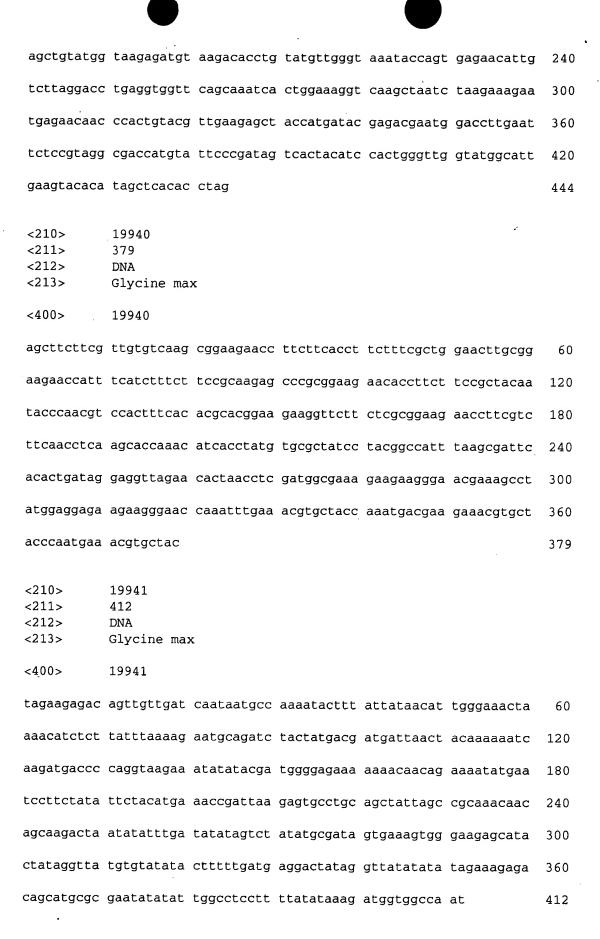
| gattggttct | ggatggaagt | cttttacaaa | ttcaaaaaat | ttggaagcta | gccaggatat | 360 |
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| tatatttgag | ttcccaaatc | caaaatctaa | ctctgttcta | ttctgcattt | gtttgtaatt | 420 |
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| <210> <211> <212> <213> | 19928 407 DNA Glycine ma | x | | | | |
| <223> <400> | unsure at a | all n locat | ions | · | | |
| agctngngat | gttctntgtt | gagcttatna | gtggaacaga | tagatgaaga | ggctgaatcg | 60 |
| ttgagaaagt | ttggataaac | ataacttgag | gaatcagaag | ctgatgatgg | aagcgaaaga | 120 |
| aagagcggaa | cttctgggag | agctgtgtgt | aattgttgac | tgtcacattc | tggtttagtg | 180 |
| attcactgtt | tctttggctt | tgctgcatat | ctgactntgc | taatttgcat | gttgctgaat | 240 |
| ggcgattcaa | gtcatgttnt | gagaatttat | gatgaagagg | cacaagcaat | gtccattcgc | 300 |
| gcgagctgga | aaatgctaat | gcagttgggg | aagccatcct | ctctacaata | catggacaaa | 360 |
| gagatcgctt | gaaggtatat | cttaactgct | gccttcactc | atcaaca | | 407 |
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| <400> | 19929 | | | | | |
| tctccctttg | ctgcagtggg | tcctctgtag | caggcaaaga | tgcttttctt | tccatgtaaa | 60 |
| tacatgcatg | tcattgaacc | atcaattaac | acacatacat | aàttgaattg | catgatgaaa | 120 |
| gattaagaga | cattcaagca | tggctatccc | acggtgcaca | tcttgtgggt | gtttggagtg | 180 |
| aacaagagcc | catgataaac | gcaggagaca | gtcttgcata | tactcttgag | atgattgttt | 240 |
| ttctgcctcc | aagacctctc | tttcacatcc | ctgccagtgc | agtgccacac | acaacaacaa | 300 |
| ttaattaatc | aattgcaatc | tcaaacactt | aaaaccctaa | cagagaaaga | gcatatgaat | 360 |
| atgaaacata | cagctacgac | gtcagagtca | caataaggaa | gatggtcctt | gccggagaag | 420 |
| aactgaccca | ccgagtccaa | gatcatccct | agettagett | caadtttccc | catttcta | 17 Ω |

| <210> <211> <212> <213> | 19930 311 DNA Glycine max | |
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| <223> <400> | unsure at all n locations | |
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| aaacttcaga | atgggccata tgctntcgaa gatatgataa agatgggtag gatgctggag | 120 |
| aggagcttct | ttttcgagcc cgaccctgct tnttcagtcg ttaacatcca ccacattaat | 180 |
| ggccacacaa | tactgatcat gagggtgacg gcagctcttt cagtgactgc atanangtat | 240 |
| gtctattctt | aaccatgtgt aagactaaat atcagctgat ggaaaataac cactcatatg | 300 |
| gtcaattgtg | a . | 311 |
| <210> <211> <212> <213> | 19931 477 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19931 | |
| cgtcgatcct | aagagtaata aaatcaaatc gctcacaaca acagtttgaa tcncctgctt | 60 |
| actactgcgc | tetgeagtgg ttatatttee eettteaaaa gagacaatgt egtgtttgta | 120 |
| ctgatcactt | tgatcacctt tgctgaacca tgagagttta aatgaagcct gtctcgataa | 180 |
| taaagaaaat | aattaaaatc aatttgctta agtaggagta aatcagagta tatacattgc | 240 |
| aattttagca | atagaaaaga ggaatatgtc tttgaagtaa aaatggtaca agatgtctta | 300 |
| ccacagattt | ctaagttcaa atacatggga tcaataatac aagatgatgg aaaaattaat | 360 |
| gaagatgtca | cgcaaaggat acaagcggga tggataaaat agagaaagga gtcaacggtt | 420 |
| atttgtaatc | gcgaagtccc taccaatatc anggcaagtt tatcgtacta ctatacg | 477 |
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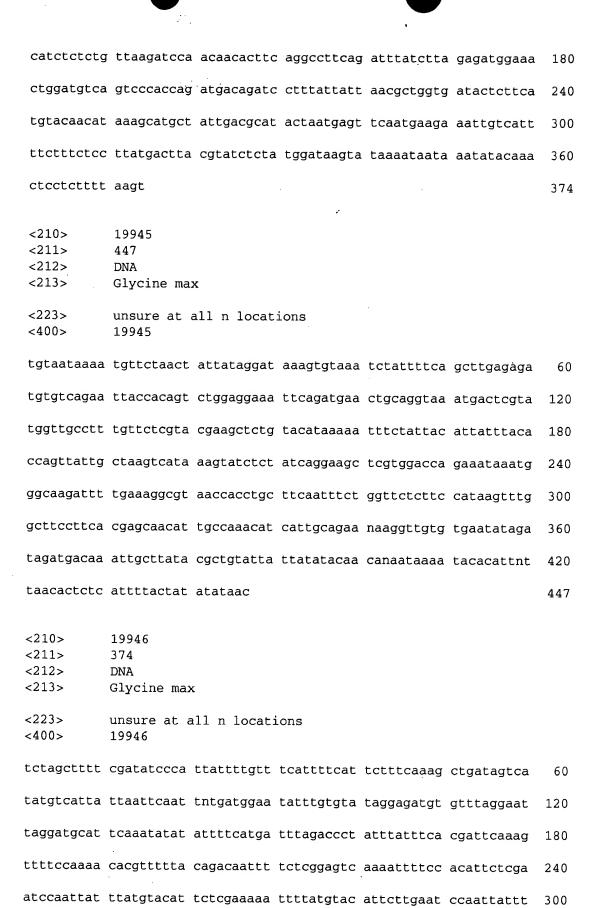




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| cttaaaagaa | ctcttgagag cctgtgtgtt agagtagacg ggtacttgga atagtttctt | 120 |
| acccttgata | gagtttacat acaacaatag ttttcactcc agtataggta tggcacctta | 180 |
| cgaggcattg | tatggtagaa gatgtaagac acctctatgt tgggtagatt ccagtgagag | 240 |
| cattgcctta | cgacctgagg tagttcacca taccattgaa aatgtcaagt tgatccaaga | 300 |
| taggatgaca | gcagcccaaa gtatgcagaa cagctactat gatcagagaa gagaggatct | 360 |
| tgaatttgct | ataagtgatc atgtatttct g | 391 |
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| agcgtgatct | tcatcgtccg cgtgtatgat attcactcca caaggtttga agtagaggag | 60 |
| accttcaacc | ctataacgca acgtgacgga caaaagtggg cagttaactt gaatggccgt | 120 |
| tattgtcaat | gcggaagtta ttctacgctt cactatccat gttcacacat tattgcagct | 180 |
| tgtggttacg | tgagcatgaa ctactaccaa tatatagatg ttgtttacat gaatgagcac | 240 |
| atcttataag | catactccgc acagtggtgg cctcttggga atgaagcggc aattcctact | 300 |
| tctgatgagg | catggacact aatccctgac ccaactaaga ttcatgcg | 348 |
| <210> <211> <212> <213> | 19939 444 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19939 | |
| caggtagagt | tcttcttacc acccatagac tgagagnctt tcctgcgctc cattcaatcc | 60 |
| ttagaggacc | tcttgagagc ctgcgtgtta aaatatgggg gtagctggga tagtttctta | 120 |
| cccttgattc | tacctacaac aatagctttc actccaatat aggcatggca ccttacaagg | 180 |



| <210> <211> <212> <213> | 19942 319 DNA Glycine ma | x | | | | |
|----------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| <400> | 19942 | | | | | |
| agcttgtcga | ttttgccatg | tttgggtgag | atagacatac | ccattctgtt | ttatggtttt | 60 |
| tgtgatgatg | tttgtgatgt | ttatatgctg | aaattgctga | tggaaatçtg | ttatagatga | 120 |
| agggtagaac | taacccaagg | ttagaaagtg | agaatgtgac | gttatgagtg | gaaaaagagt | 180 |
| gagactctga | gagttggaag | gctaattctg | aattctgtgg | taaatggatg | ttagagtgag | 240 |
| ttaatactag | ctagaaatgt | catttataac | atgtgagaaa | ggttaagctg | agctatagag | 300 |
| aaaaacaaat | gaccaaagt | | | | | 319 |
| <210> <211> <212> <213> | 19943 412 DNA Glycine max | × | | | | |
| <400> | 19943 | | | | | |
| ctcaactgaa | tttacaacat | tccaattgat | ttcaaaatgg | tgtattttat | tacaatgata | 60 |
| tggtaatcga | ttaccagtgt | gtttgaatgt | tgaaattcat | attcaattgc | gaagagtcac | 120 |
| atcctttcac | ataagtgctg | tatgtaatcg | attacaatga | tttggcaatc | gattaccagg | 180 |
| gatgtgtttt | gaatacaaat | cactagatgt | aactcttcca | atggttctca | agtctctcta | 240 |
| aaggctataa | ctcatctatt | ggccttcttg | acctgacttg | acgagtctat | ataaccaaga | 300 |
| ccttaacttg | cattgtacac | acattgatta | caatcttata | tatcctttga | atctctttga | 360 |
| acctcctctt | gaatgtcttc | ttatcttcct | ttgccaaagc | tttctaaagt | tt | 412 |
| <210> <211> <212> <213> | 19944 374 DNA Glycine max | | | | • | |
| <223> <400> | unsure at a 19944 | ıll n locati | ons | | | |
| agcttctaat | gtataacaaa | gtgactctat | nttcagggtg | aatcattacg | ctgcttgtcc | 60 |
| tgaaatgact | ctgaatgatc | acaacttgat | tgggtttgga | gaacacacag | acccacaaat | 120 |



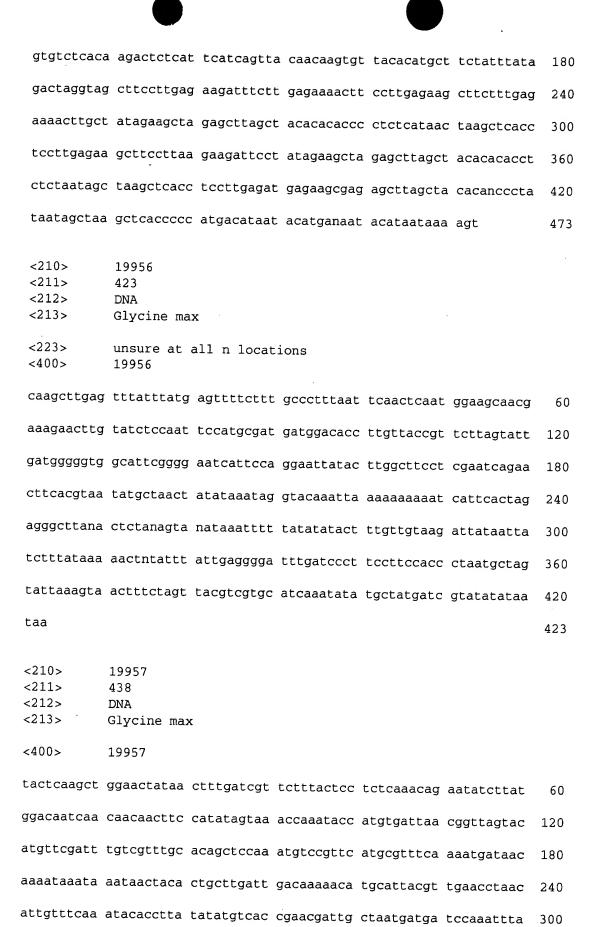
| atgcacattc | tcgaagctga | gtcatatctc | gaatcçaatt | atttatgtac | attctcgaag | 360 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| ctgagtcata | tctc | · | | · | | 374 |
| <210> <211> <212> <213> | 19947 434 DNA Glycine max | × | | | | |
| <223> <400> | unsure at a | all n locat | ions | | | |
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| gctcataggt | taagaacact | tcatgataac | aaagatgatg | atctcaagaa | tcaaagaatg | 120 |
| agttcaagat | gttcaagatt | gaatcaagaa | catttcaagg | ttcaagagga | aatttgattt | 180 |
| caagaatcaa | gaatcaagat | tcaaggttca | agcttccaag | aatcaagatc | aagattcaag | 240 |
| actcaagatt | caagaatcaa | gagaagactt | aatcaagata | agtatgaaaa | agttctttca | 300 |
| aaaactgagt | agcacatgga | tttttctcaa | aacctgttta | ccaaaggggc | gtattaccta | 360 |
| accggtaatc | agaanatgca | gaaccataga | ccccaggccc | atacacccca | agcctagtaa | 420 |
| ggtcactgta | taaa | | | | | 434 |
| <210> <211> <212> <213> | 19948 435 DNA Glycine max | ¢ . | | | | |
| <223> <400> | unsure at a | all n locati | ions | | | |
| agcttttcat | ttaccttcac | aataagggaa | aaatgttatt | agaactaatt | aataaataca | 60 |
| aaaaagaact | aaaatgacag | ccataagtat | gtacctttga | ttntttcttc | aaattataat | 120 |
| tatggcgaag | tcaatccccc | tgcaaatgag | cgcttcaata | gattcttcat | ttagtttaga | 180 |
| acaatactca | tcaattaccc | tacccttagc | actaaatgta | gactctgaaa | ctactgttga | 240 |
| tattggaata | gctagtatgt | cacccgccat | ctttgataaa | accttgtatt | ttaggctatt | 300 |
| gttcctccac | cactctaaca | cactanaata | agagttacta | gtttcaggaa | tataaacatt | 360 |
| ctccttaaga | taatcctcta | attctgagtt | cactggaggg | gtggcttcat | ttgcacgcac | 420 |
| aatgttcatt | attto | | | | | 435 |

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| acaanaaatg | aacaagattg ataagccaac atatctagta taaacaaaat caccaccaca | 120 |
| caatttgtat | gctaattata aagaattcta attcctaagg tacacaccta acaaaggaac | 180 |
| acatcaattc | tacaacaaac tcgtatcaga acaccaatta gttcatcaaa cacactcaat | 240 |
| ccgtaattaa | acatgaaaac ataattaaac ttcataaaca ccccaaaata acccaaaaat | 300 |
| tgatcctcta | aggateceta cacatgitea itetaatiee caagegigag taacteatee | 360 |
| cttacgtcga | tgtagtcgct cacatattct ccgctagtaa ttgtggcgtc tctggtgctc | 420 |
| tctagagctc | ctactctggt tgttct | 446 |
| <210> <211> <212> <213> | 19950 ° 450 DNA Glycine max | |
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| ttaactcatc | gccttaaatg gtcttatagt catgtgattg tacaattcat agatcacaac | 120 |
| tcanagcaca | taacatctca atgcatatat atcacataca ttcggtctca atcacgatga | 180 |
| tataatccca | gagtaacatg ctatcacacc tcatgaatca tatgcacttt aattatgaac | 240 |
| tatacaatac | acacaattac tcatttgttt tcaaaatcat tttaactcct cgcacctcan | 300 |
| agtgattcaa | ctcatcgggt tcccatagtg gatcccatca gaatactcgt cgtgcaaaaa | 360 |
| ctcgctgctc | ttaaagggtc ttacaattgt gtgattgcac agttcatagt tcacaactca | 420 |
| atatatacaa | naatgatgta gctccatgtg | 450 |
| <210> <211> | 19951 450 | |

| <212> <213> | DNA Glycine max | × | | | | |
|---|--|--|---|--|--|--|
| <223> <400> | unsure at a | all n locat | ions | | | |
| tgagatgagg | aagtgtagaa | gggtgaaact | tcctgctntt | attcgttgac | cacatagtgg | 60 |
| tacctggaca | tatgtcgcgg | tggtcaggag | accttgtgga | cgtcaggtgg | ggtgctactg | 120 |
| cccaaaacca | agcttgacca | atcccgaccc | aacccgggca | tagtcagtca | gtgataacct | 180 |
| gtgatgtacc | taaacaggcg | agctcctggc | agtcaataga | taaaaggaac | aaagaccaca | 240 |
| aagcaaggag | gcttgtggtg | gctggccagc | tgtgaatttt | gtgtgatata | tgggttgtgg | 300 |
| cctctggtaa | tcgattacca | agggtgggta | atcgattaca | aggcttgaaa | atgaagacag | 360 |
| gaggctaaga | tggtctctgg | taatcgaata | ccacggggtg | taatcgatta | ccangcttga | 420 |
| aaacgaggtc | aggaagctaa | gggagcttct | | <u>.</u> 34 | | 450 |
| <210> <211> <212> <213> | 19952 442 DNA Glycine max | c | | | | |
| | _ | | | | | |
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| <400> | unsure at a | all n locat: ggcaagttca | acatgctttc | | | 60 120 |
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| <400> agcttaactc actatcatga acgaaaatca acacgcactt caaaggttga | unsure at a 19952 atccaatcat agcaganaac agagagaaac cacgactccg | ggcaagttca tagcaaaact aagtccaccc aanatgccct gagcttcaat | acatgettte acceateata aaacetgaaa cetttegega ggagaatgga | tctcccanaa tttcgaagtc tttggagcag ggagaacgaa | ccccataccc tcacacgtag aaatgggcac naagcaacgt | 120 180 240 |
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| <pre><400> agcttaactc actatcatga acgaaaatca acacgcactt caaaggttga gaggaagagg cttttggttt</pre> | unsure at a 19952 atccaatcat agcaganaac agagagaaac cacgactccg agctntgttg gagagagang | ggcaagttca tagcaaaact aagtccaccc aanatgccct gagcttcaat ctttgaaaat agcttttcct | acatgctttc acccatcata aaacctgaaa cctttcgcga ggagaatgga gtggggctga | tctcccanaa tttcgaagtc tttggagcag ggagaacgaa gtgaggagag | ccccataccc tcacacgtag aaatgggcac naagcaacgt agagangttg | 120 180 240 300 360 |
| <pre><400> agcttaactc actatcatga acgaaaatca acacgcactt caaaggttga gaggaagagg cttttggttt</pre> | unsure at a 19952 atccaatcat agcaganaac agagagaaac cacgactccg agctntgttg gagagagang anaaannaaa | ggcaagttca tagcaaaact aagtccaccc aanatgccct gagcttcaat ctttgaaaat agctttcct | acatgctttc acccatcata aaacctgaaa cctttcgcga ggagaatgga gtggggctga | tctcccanaa tttcgaagtc tttggagcag ggagaacgaa gtgaggagag | ccccataccc tcacacgtag aaatgggcac naagcaacgt agagangttg | 120 180 240 300 360 420 |

| tggtttgagg | j tacttacccg | ttgaagactg | aagaaaacga | agaatggact | atgaatcttg | 60 |
|----------------------------------|---|--------------|------------|------------|--------------|-----|
| aagaacggto | gagaatcttc | gcgtaattac | tcacggaaac | gttacggaag | g cgcctcggct | 120 |
| tggattttct | tcatggaact | aattttcctc | agcaatttcg | agagagagag | , aagtgcctaa | 180 |
| ggggttgaad | ccttttcttc | ttcacttctc | cccctattta | tagcaaaata | ggggttgtat | 240 |
| atcctcaaat | aataatcccc | ggacaaaatt | agggtatgac | agttgcccct | ctttacttac | 300 |
| ctctcatcgg | agataagagg | aaagcaaaga | taggacactg | atttcgtccg | tcctgccctt | 360 |
| ttccgtgatg | acgactctcg | tctctactcc | tttcttttt | cttctgcaca | aaacaaaata | 420 |
| catacaacaa | ccagaacaac | gaatataata | cacatataca | catatacaca | tact | 474 |
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| <400> | 19954 | | | | | |
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| cctgtcgcaa | gagtctgtgg t | ttatggctc | tctgctgacc | accatacata | ccttttgcct | 120 |
| ttcatggcgc | aacctggaac a | aatagagcaa | cctgaagctt | atgttgcaaa | catttacaat | 180 |
| agacactctc | cacctcagct a | acaaaatcta | ccgcagcaga | acaattatga | cctctccagg | 240 |
| gaccattcgt | tgttgtgatg c | cgacccctat | ttgaccactt | ggaggtgctt | ggcacccatc | 300 |
| gctatgcgat | ttgtgatagt t | cctgacatg | ccgcgaatct | aaagaaacac | tgtgcacaaa | 360 |
| acccgtagat | tcccgcatgt g | gcgagatata (| aagagaagtg | gtgtgtactc | tcgaagagtt | 420 |
| tgcacttacg | ggat | | | | | 434 |
| | 19955 473 DNA Glycine max unsure at al 19955 | l n locatio | ons | | | |
| gacctataga | tactaagctg n | agaggatgc t | taatggagg | aaaagaaaga | gggagagaaa | 60 |

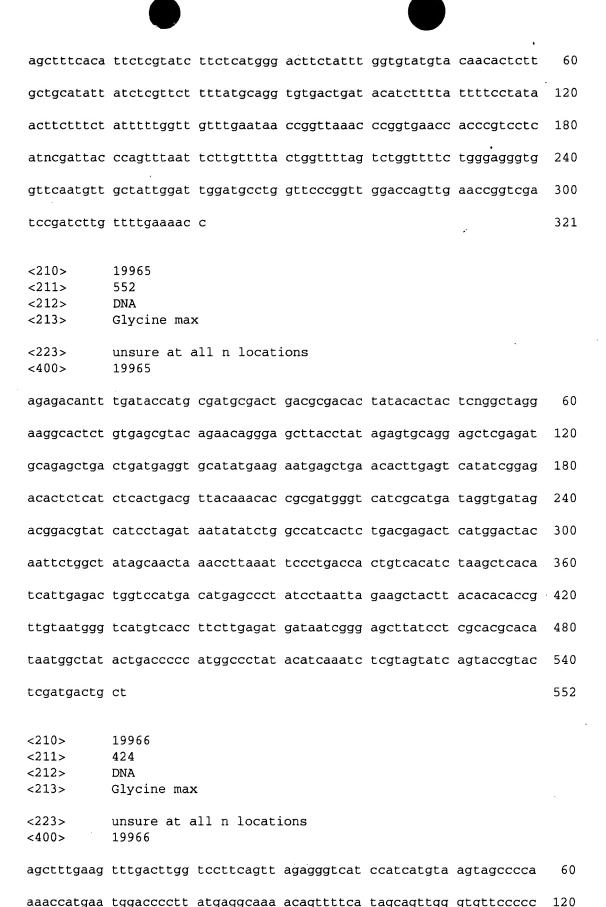
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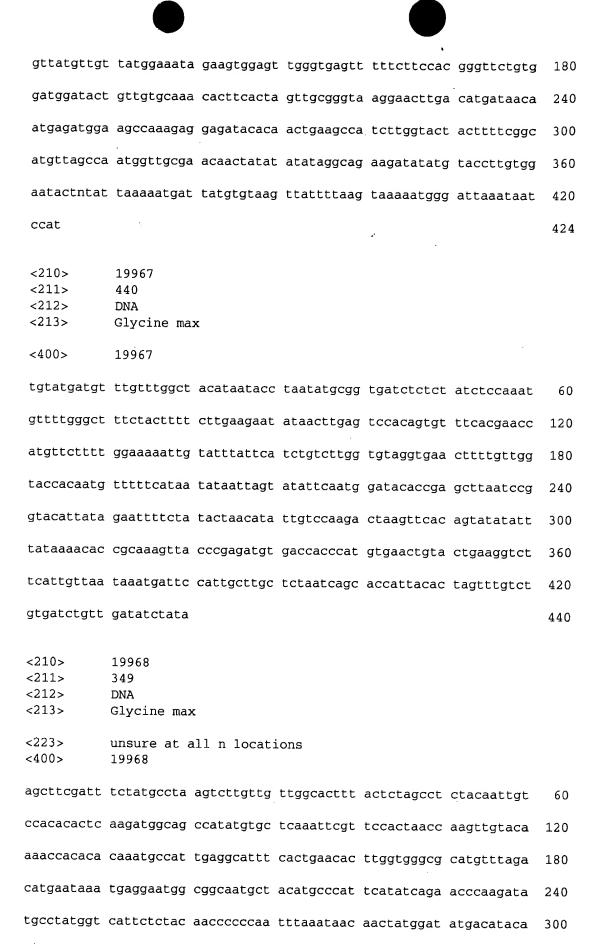


| agattataag | aaataacttt | gtttatagta | ccggaaaaat | tgagtatgaa | actttgacaa | 360 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| acatattacg | aaaataatac | tatatagaaa | caataaaata | atatatatat | ttataaatac | 420 |
| tctaacattt | atcttcat | | | | | 438 |
| <210> <211> <212> <213> | 19958 392 DNA Glycine mas | × | | | | |
| <223> <400> | unsure at a | all n locat | ions | | | |
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| aatcggtacc | tgtcgcaaga | gtctgtggtt | tatgctcctc | tgctgaccac | catacatacc | 120 |
| tttgcccttc | catgcagcaa | cctggaacaa | ttgagcagcc | tgaagcttat | gttgcaaaca | 180 |
| tttacaatag | acctcctcaa | cctcagcaac | aaaatcaacc | acagcagaac | aattatgacc | 240 |
| tctccagggg | accatccgtt | gttgggatgc | gaccctcatt | ngaccacttc | gaggtacttg | 300 |
| gcacccatcg | ttaggcaatt | tgtgaagttc | catgacgtgc | cggaagtcga | aagaaagcat | 360 |
| tgtagcacga | tccgtgaagt | tccgcgacat | gc | | | 392 |
| <210><211><212><213> | 19959 476 DNA Glycine max | ζ | | | | |
| <223> <400> | unsure at a 19959 | all n locati | lons | | | |
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| aaaattgaag | gaagaaaaag | ggagagaatt | tgaactttga | gttgtgtctc | acaagactct | 120 |
| cattcatcag | ttacaacaag | tgttacacat | gcttctattt | atagactagg | tagcttcctt | 180 |
| gagaagattt | cttgagaaaa | cttccttgag | aagcttcttt | gagaaaactt | cctatagaag | 240 |
| ctagagctta | gctacacaca | cccctctcat | aactaagctc | acctccttga | gaagcttcct | 300 |
| taagaagatt | cctatagaag | ctagagctta | gctacacaca | cctctctaat | agctaagctc | 360 |
| acctccttga | gatgagaagc | gagagcttag | ctacacaccc | ctataatagc | taagctcacc | 420 |
| cncatgacaa | aatacatgaa | aataccaaaa | anaagtccct | actacaatga | ctactc | 476 |

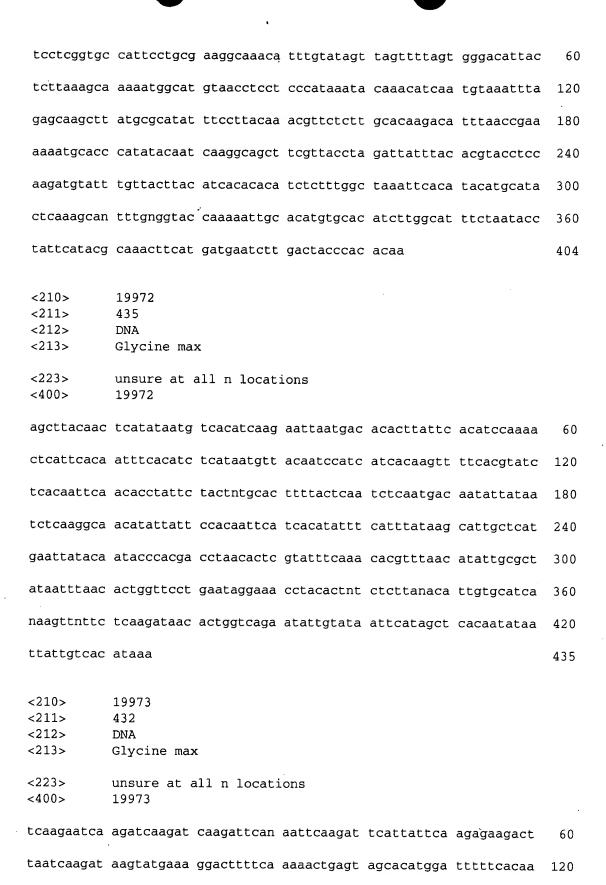
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| aaaagacata | atattaaaaa t | ttcaagcaca | tgtaataacg | taaaataatg | tgggaaaata | 120 |
| catatttaca | actcacctgg t | tcgatcgtta | gtttcttgaa | actgtaaaat | gagatgacct | 180 |
| ataagaagct | agacattaga (| cgtgacactg | gttgagtcag | aagtgtatat | ctggccctat | 240 |
| ttattataat | tnttatctgc (| cccacatga | aatnttaatg | gtatttaggt | accgtagata | 300 |
| tacatattac | ctgtaatata t | tatgcaatct | ctcaatactt | atatngctgt | tattaaccca | 360 |
| tttctttgca | gatttgaaag a | atataactct | cttccaccca | ct | | 402 |
| <210> <211> <212> <213> | 19961 470 DNA Glycine max | | | | | |
| <223> <400> | unsure at al 19961 | l n locati | ons | | | |
| tgaccttntc | ccatggactc a | aggttcaac | gcaacaggct | tatcttcctt | gngtgaatca | 60 |
| aatgccgtct | acagtatgat c | catcactcta | tattaatttc | aggtaaagaa | aaatggtgcc | 120 |
| atttggatcg | cataaacagc a | actacctta | tạccaagtga | gtgtttgatt | gcaagtattc | 180 |
| cctagtggac | tccattgaat a | gaactgtta | cttttttctt | cgtatatttc | taattgctct | 240 |
| cccagtaatc | caacctgtaa t | aacgtacgt | acatgacacc | aatgtcctac | tgtccaatta | 300 |
| aatctataca | acctaacaga t | ttgcagtaa | ttaaatgtgc | ttgagaataa | tatgcgttag | 360 |
| ttccacgagg | aatacacata t | ataatattt | aattatatat | aaactcatca | taatacaaga | 420 |
| gcaaagcgga | cctgatatcc n | catgtagaa | ttggtcatat | cgtaggactc | | 470 |
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| <223> <400> | unsure at all n locations 19962 | |
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| agcttcatg | c tgtttaattg ctccaggttg ctgcatggaa gggcaaaggt ctgtatggtg | 60 |
| gtcagcaga | g gagcacaaac cacaaaccct tgcgacaggt acagatttct gattcaaggc | 120 |
| cagctgggt | accaagttaa ccaatgcatc cagtttgcct tcaagcttct tagtttcaga | 180 |
| tgatgcagai | gggtttgtag ctacctcatg cactcctcta atgactatgg catcatttct | 240 |
| ggcgctaaad | tgctgggagt tggaggccat cttctcaatt aaatttctgg cttcagcagg | 300 |
| agtcatgtct | ccaagggete caccactgge ageatetate atacttetet ccatattact | 360 |
| gagteettea | taaaaatatt ggagaagaag ctgttctgaa atctgatggt ggnggcaact | 420 |
| ggcacatagt | ttc | 433 |
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| ataaactntt | agaagcaacc aatttgtggt actcagagca gagagatatc tgttggtttt | 120 |
| caccaataga | ggaagatgtt gctatgtggc tttttgtctc gactacctct gaccttgatt | 180 |
| cttgccataa | tcatgtttca acttcaagta gttatgatat acatacagct agaagttgga | 240 |
| ggcttgctct | tttgatgaaa aatagcataa tttttggaag tcccttggat ctaaggtgct | 300 |
| atgaagtttt | ttctgaattt gaatatagtt cgtcatgtgt tgcttaattc tgttaagatc | 360 |
| tttactggtt | ttcacttcac atctagattt aattgggaag attgtaagga accattagtt | 420 |
| atatatcatt | ttgagtgata tggaaggata taac | 454 |
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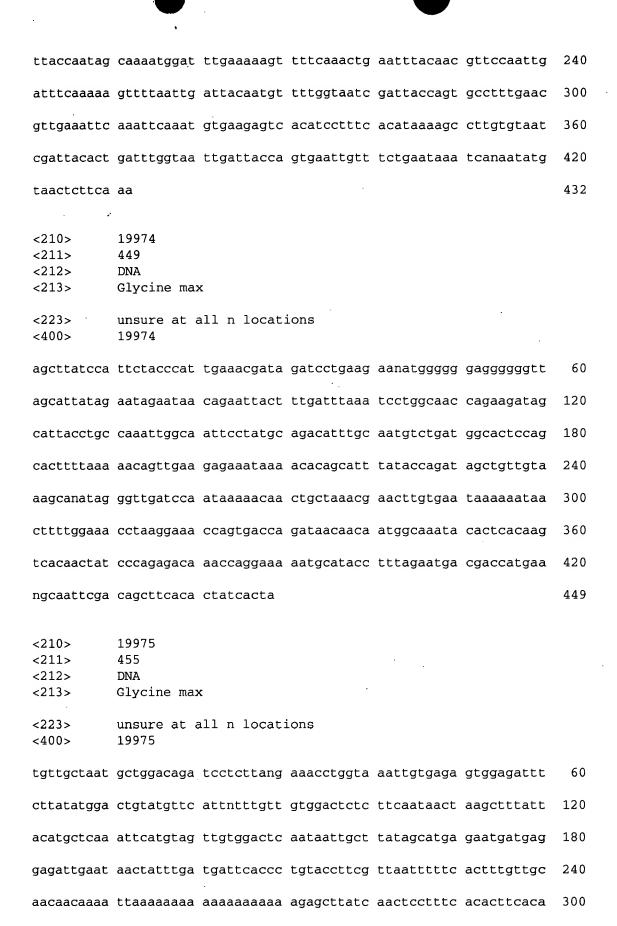




| ttgtctcatg | gattttgcan | acatagacaa | ttaaagcact | aaaacacat | | 349 |
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| atgttcatca | tggatttatg | ggatgtaatg | aaaaaatatt | ttttggcatt | tcttgatgaa | 120 |
| ttacacataa | atgggagact | agataaagtg | gtcagtagat | cttttattgt | tctgttacct | 180 |
| gaaaaaagaa | aacctaatgt | ctatggggga | taatatatcc | ctgatagggt | gtttgtataa | 240 |
| aatgttggca | cagatgtttg | ctaataagtt | aaaatgggtt | attgatgatg | ttatttccac | 300 |
| aacccaatct | acttttatat | cagggaggaa | aatgctggat | cgggtactca | ttgctattga | 360 |
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| ttgggcaaaa | ttggatgagg | gaaagtgtga | tttcgaaaat | ctgcacttta | tgcagaattt | 120 |
| tgctgtcaaa | taggtgcaac | agaattttgg | ctttgtgcag | aaagtgttgt | gtaattgctg | 180 |
| gctgtggaaa | gagtagtata | gattgtgttc | tggacgtttt | ctagcagatt | ccaacggtca | 240 |
| taatgtagat | ttatgtgcta | gagacttcca | gtaaaatttt | cgagtcgatc | caactgttaa | 300 |
| cgaattggaa | cgaagagaat | attactggcg | tatttgaatg | ttgaaagctg | tgata | 355 |
| <212> <213> | 19971 404 DNA Glycine max | | | | | |
| | unsure at a 19971 | 11 n locati | ons | | | |



aacatgttta ccaaagagtt tttactctct ggtaatcgat taccagatta ttgtaattga 180



| ggaaaaagag | cctgaaactt | gtgtaccaca | aacaagttct | agtcattccc | aagttttcat | 360 |
|-------------------------|------------------------------------|-------------------|------------|------------|------------|-----|
| tgttatttgt | tagtcctcgt | acacaaaact | tgaattcttt | gaacttgttt | gaacaaatac | 420 |
| ttattagtac | ttanatttcc | tccatttcta | aatat | | | 455 |
| <210> <211> <212> <213> | 19976 408 DNA Glycine max | ĸ | | | | |
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| agctintgcg | ttcgatttct | tagtgcaatt | ctccattctc | aacctttttc | ggagccccat | 60 |
| ggattgagtt | ttcgttcatg | cgtactccac | cttcgagtat | ggagccatgc | gtagtgattg | 120 |
| cttagttcaa | ttctccattc | tcaacccctt | tttcgcagcc | ccatgaattg | cgatttggtt | 180 |
| catgtgtcct | ccaccttcga | gtctggagcc | atgcgtagtg | attgcttagt | gcaattctcc | 240 |
| attctccacc | ctttgtcgga | gcccatgaat | tgcgtattcg | ttcatgtgtc | ctccaccttc | 300 |
| gagtttgaag | ctctgcgtag | tgatttctta | gtgcaattct | ccattctcaa | gctttatcgg | 360 |
| agccccatga | attgagttat | cgttcatgcc | tcctccacct | tcgagttt | | 408 |
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| tatattcacc | aagaaaacat | tatagtggaa | ggacattgta | gtcttgtgat | tcaaaagatc | 120 |
| cttccaccca | agcataaaga | ccttgggagt | ataactattt | cttgttcaat | tggagaagtc | 180 |
| actatgggaa | aagctcttat | tgacctgnga | gccagtataa | atttaatgtt | gctctccatg | 240 |
| tgtagaatgt | tgggagcgtt | agagatcatg | cccactagaa | tgactctaca | attggctgac | 300 |
| cgctccatta | ccagaccata | tggagtaatt | gcagatgtgc | tggtcaaagg | gaaacatctc | 360 |
| atcttcccgg | tagacttcgt | ggtattggat | atttgtgaat | atactgacat | tcctgtaata | 420 |
| ttggga | | | | | | 426 |

| <210> <211> <212> <213> | 19978 376 DNA Glycine max | |
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| <223> <400> | unsure at all n locations 19978 | |
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| tggctttccc | tttttggtga gctttgggaa gaacctggac agacatgcta gcctttcatt | 120 |
| cagetteage | ttctacactt cttggatgtt ggttgggttg cgcatgatca gtatggtagt | 180 |
| gcatttgttg | nggttggctt caatccccta gtgagtgatc atgaagtcga ggaacatgcc | 240 |
| tctgtctacc | caaacagtac atttatcatg gttgaggcac atgtcatatt agtgaagttt | 300 |
| ccacaagact | tettecaggt cagteacatg ttgggetatg eteegagaca tgtetatgat | 360 |
| gtcgtgcaca | tatacc | 376 |
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| tgacagcccc | ctatccattg aatgtctcag gtgttttact gctcttttaa ctgcaaagat | 120 |
| tatttatatt | acagtaggac aagtaccata ggcactgcgt gatacatacg ataatataat | 180 |
| cctgtaagaa | cagaaaaaac aacccgacat gaagccacat ccctacccta | 240 |
| attaactaaa | aactagaggc attgttggaa tcccatgtgc tggccttcat aagtctatcc | 300 |
| cataacaatc | tccaggcata agataaagct ctaggaggga ttttaatctc ccatagttga | 360 |
| tggaagccca | ggtgctggcc ttcataaagc tgctcagctt taatgac | 407 |
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| <223> | unsure at all n locations | |

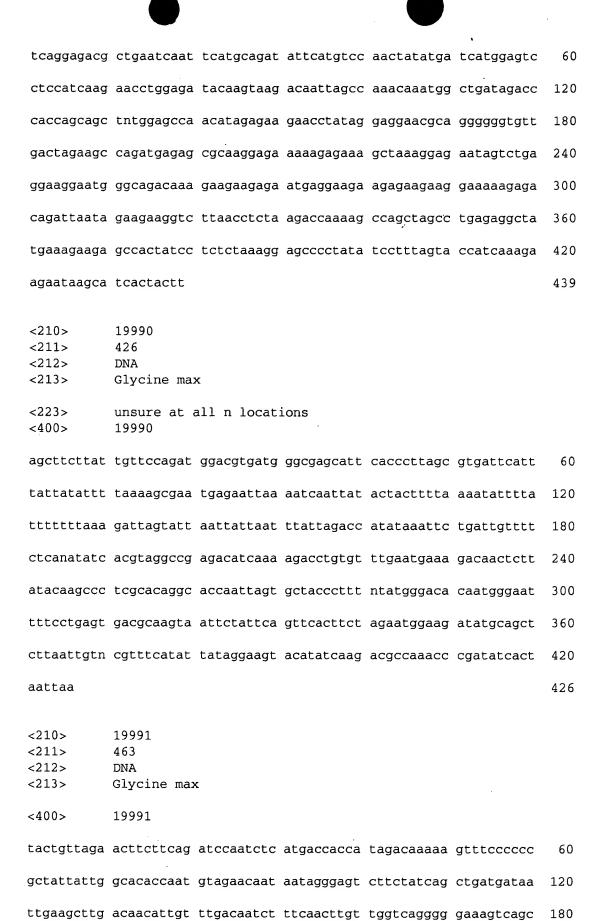
| agctttgaag | aaacaattac | aagctatcta | cgtgaattat | tctttcatac | tttntgtaaa | 60 |
|----------------------------------|------------------------------------|---------------|------------|-----------------|------------|-----|
| tacttttaaa | gatacaaagc | tctcaaaaca | ccttgtatac | tttgagagaa | aacactaaaa | 120 |
| gtgttgagtg | ttatatctat | ttgtaagatg | attatatatt | ttagttagtg | tgaaaacttc | 180 |
| taacaaatct | tgttgatttg | tttagagcca | gatgtggctt | ggtaggacaa | ggagtactgg | 240 |
| gttttaaatc | aaggttgggg | gtagagcttg | caagtgaaag | agtcagaagt | ggcaataagt | 300 |
| aatacttgta | actntgataa | gttagtggaa | acttggtggg | tgctaagaat | tggacttagt | 360 |
| ctcgaggttg | agacaaacta | gtataaattc | cttgtgtggt | ttttgttgtt | aactaattca | 420 |
| taccctaatt | tc | | | | | 432 |
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| ttttggttat | tggctctggt | agagccagta | gaacctcttt | tccttttgta | ctggatcttc | 120 |
| ctcctctgcc | ggaggaacta | gacaccatag | gaggcatcgg | gaattttctg | taaaaattca | 180 |
| cgggtgagat | agtcaggtag | agagttggaa | gagcccttga | tatattctat | attaaaatca | 240 |
| aagacactta | aaattgcttg | ccatcttgca | aaaatctgtt | ttgaggcaag | gttttttaca | 300 |
| tccttctgta | naatgtcttt | ggctgatttg | cagtcaaccc | ttactaaaaa | tttttgattt | 360 |
| aataaatcag | attgaaattt | ggaaatgcac | aaaacaattg | ctaaaacttc | ttttttgaca | 420 |
| gttgaatact | ttaattgtgc | aggattccag | tgttntgaag | tatatgcaat | ga | 472 |
| <210> <211> <212> <213> | 19982 420 DNA Glycine max | | | | | |
| <223> <400> | unsure at a 19982 | ll n locati | ons | | | |
| agcintgact | tatccaatga | anaaanntan | agastaatsa | + ~+ + + ~+ + - | | |

ctatttttt agggggagtt tgatttggtt tgaaagaagg agtaagataa gaagtgagag 120

| atgattcact | ctgattgcca | tttgtttgag | agaggtgcta | gagaattggt | cgcactatta | 180 |
|----------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| aatgggacaa | agggaaatcc | ttggacattt | tatcatagtc | acttggtgag | ttntaatcta | 240 |
| acacactcct | acattttata | aaacatgtat | taatgatggc | tcaaagttac | gtacaagtct | 300 |
| gtggctgggt | gctatttaat | cgttggataa | ctctaaagtg | cttttgacta | cttacaataa | 360 |
| tttaaggata | ctcccatacc | tttgtgaata | attactcatt | tgacattgtt | tacacatgac | 420 |
| | 19983 435 DNA Glycine max | ς. | | | | |
| | | gaacatgtga | ttagaattat | gaatgttatg | ctagttttt | 60 |
| | | | | | | 120 |
| | | | | tgagagtgtg | | |
| ttgagagaaa | acggctatca | ttaagtactg | acttttgcat | gaatctctta | attatggact | 180 |
| gaatgcatga | atttgaggat | gatgaaggcc | atgttttgat | tgtgatagcc | acttagccaa | 240 |
| aaagctgacc | atgtgcatga | atgatttatc | ccttgcaccc | agttttgagc | tgaatgaatg | 300 |
| cttgattgat | tgaaccttga | gcctatacag | ttttatcttc | tgctactttg | tcttacgttg | 360 |
| taggagagca | tcatccacag | aaaagcttag | ttcaaggcag | atttgtccca | aatttgggga | 420 |
| gttatatgtc | aaaat | | | | | 435 |
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| atgagtttat | cagcaactca | ggattcaaca | gatgtgacat | gaaccattgc | tgctatgtta | 120 |
| agaaatatac | taatagttat | gttatcctta | tcgtgtatgt | tgatgacatg | ttgattgcag | 180 |
| gatctagtat | gacagaaatt | aacaggttga | agcaacagtt | ggcagaaaac | tttgaaatga | 240 |
| aggatcttgg | tccagctaaa | caaatccttg | gtatgagaat | tcttagaaac | agatcagaaa | 300 |

| gaattttgäa | gttgtctcat | gagaaatata | tacacaaagt | tgcttgacaa | gttttacctt | 360 |
|----------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| gaagattcta | agaccaggaa | tacccctttg | ggatctcatt | agaagtttca | aag | 413 |
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| cggcaaccgt | ttggtactag | aaaacctact | tccatgcttt | atttgtaaaa | acatcttata | 60 |
| taatcatgca | tacaagtgac | aacttattca | tattctcctc | tctaattttt | ctctggcata | 120 |
| aaaagcgaac | aataaattca | aaagcatata | aggagataat | atgatatgta | taaaaatatg | 180 |
| ccccttttct | tcccagagac | aatttcatct | cattttataa | caacaaagtt | gcttagtact | 240 |
| aaatattatg | atcactttgg | tgtgactact | cagggaatca | ccacatatga | ttacatcata | 300 |
| gccttgaggg | agcacgagca | ggagcaacaa | ggcattggag | gtcagcagag | tccccatatg | 360 |
| tcgcctgtta | gctcccttac | tggaatgagc | agtgcaagc | | | 399 |
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| <223> <400> | unsure at a 19986 | ıll n locati | ions | | | |
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| ggggttttag | aggctcagtt | ttgtgttgga | catattgaca | atgatgatgc | aaatgaggtt | 120 |
| tcaccagctt | ctaaaaggaa | aaaagtcaca | gctaacccaa | atttcacaaa | atgtcaatcg | 180 |
| gaattgagtg | ctgtgattgt | gccttccacc | acatctgaag | ctcctatgag | tttcagtgat | 240 |
| aaccaagaac | atcagagaga | agttgctttt | gaaagcatgg | gtatgattat | tctgtctagt | 300 |
| gctcagtcaa | tgccttattc | agaggatatt | actaanatgc | ctgagaatgt | tttggctggn | 360 |
| ggatcttttg | agtctattga | tgcanataag | gaaaccatga | gttctgagca | tttggaattg | 420 |
| ggcattcgga | tatag | | | | | 435 |
| <210> <211> | 19987 453 | | | | | |

| <212> <213> | DNA Glycine max | • |
|-------------------------------------|---|-----|
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| ccttccatag | tcatcttgca nacacatget egteeetgtt atgtttttgt teatatggaa | 120 |
| gactcaaata | tacattgcac totatccaag gottootoca ogatotottt acactottoc | 180 |
| agccatggag | aaactcttca cctctcttaa taatgacttt ctcattaagg acctccttat | 240 |
| tctcctaggt | tttatatatc attactcctt catatgcttc cataacaaca tggccatact | 300 |
| ttgttgattt | gctatcaacc ttgaaagcat gttgaagaac agtngattac atgaatctgc | 360 |
| ttgaaggatc | gagccaacga taagcaaggc ctatagaaaa tattaacagt cgattacttt | 420 |
| aaatatatat | gttggactgt ttgctaacat ata | 453 |
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| gatgcaatat | ctgtggtgag gcccatgagt caagcatgtg catggtccaa gatgatgtat | 120 |
| ccaaagaagt | caactgtatg atcattccac accatcaagg gttccatcaa ggacgacctc | 180 |
| caggatacaa | tcagggggga aaattctctc agggccaagg ttggagatcc caccccggga | 240 |
| atagcttcaa | caaaaatcat ggagtttcat ctaatcagcc tcccaatcaa tggcctgatt | 300 |
| tatatgagat | aaccactaag ctagaagaca ctctgaatca gtttatgtag gtttctctat | 360 |
| canatcataa | gagcactgag tctgccatca gaaatct | 397 |
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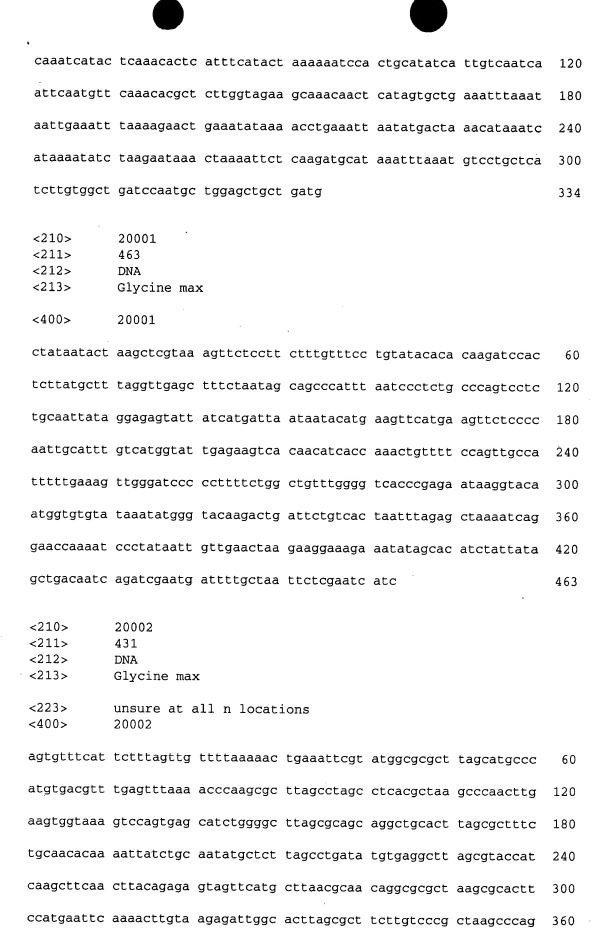


| agttgacaaa | tcccaaaaca | ttatatcacc | atctacataa | cccacaacaa | ccaccgatcc | 240 |
|----------------------------------|------------------------------------|------------|------------|------------|--------------------|-----|
| atcattagat | gcccaagata | cagagcttat | ctccttatcc | tcctcttcat | ggtctaattt | 300 |
| atcatcagaa | agctgaaccc | tagagtcatt | tggataacta | gtcactattt | ttctcttcaa | 360 |
| tttgatgtcc | ttgtggcctc | taatgagaac | aattcgatct | tcagaagcat | cccagagtac | 420 |
| catcaaacca | ttttcgtatg | caattagcag | tctgcaaaat | gac | | 463 |
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| taagcttaaa | gaatacagac | aacacaaata | tcatcgtgaa | tgatgagaag | aatatccaaa | 120 |
| caaccagccc | tctcttgcca | gttgatgggt | ctttaatatt | tgttgaatcc | ggctagcttc | 180 |
| aaccaattct | aacaccgacc | acgcaaccac | aaagtgcatc | ctttggttgt | actcaaaaca | 240 |
| gggagacttg | gcacatgtga | acctctggaa | caagtttcat | tcagcacttc | tggttttgct | 300 |
| gatagcatgt | gccaacaaca | aggggattga | tataccatag | cctaagaaat | ggagtactgg | 360 |
| gttcatctca | tatagcttta | gaagagttct | gcaattctgt | aatggaatag | tagtaaatca | 420 |
| caaaatg | | • | | | | 427 |
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| cttcatatct | cccaaaaccc | catacccacg | aaatttaaga | gagaaagaag | tccacccata | 120 |
| cctgaatttt | cgaagtccca | ctcgtagcca | cgcacttcac | gaccccgaat | atgccctcct | 180 |
| ttcgcgattt | ggagcagaaa | tgatggccaa | aggttggagc | tttgtgtgga | tcttcaatgg | 240 |
| agaatgaaga | agaagagaat | ggcaacgtga | gggagagaga | gagctgtctg | aaattttgt g | 300 |
| gcgctgagtg | aagagagaga | gagttgctcc | ttgggtttaa | atgacagggt | cttctctatt | 360 |

| tttctattat | tctattcaag | ctatgccaca | tgtcttcatt | tgagtggagc | aagaatgccc | 420 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| actttccctt | cttaattgtg | actcatact | | | | 449 |
| <210> <211> <212> <213> | 19994 465 DNA Glycine max | κ ` | | | | |
| <400> | 19994 | * | | | | |
| tctacttatg | tggcagggcg | ggcttccttc | accttcttgt | ctcctacgct | aactttgacc | 60 |
| actgttcttc | cttcccgcga | tgcttctttt | cacgtccgcc | tgagtgggct | tatagcctaa | 120 |
| accatacttc | ccacgatttc | cttgggtatt | tatcaggcta | gttatgccgc | cgttgcttct | 180 |
| gcctaaaccc | atcctgagtt | cataaccgtt | ccccaacata | actcgggcca | tcattaccgc | 240 |
| tgcatcggac | atacaaggct | gcccatagag | ggagtccacg | gatgaaatgc | tgaccacctc | 300 |
| gaaagactgg | agagcagttt | ctaacgattc | ttccgcggct | tccacattag | gcatggagga | 360 |
| tgggcagctc | accaagatat | tttactcgtc | tgcacgatga | ccaagtgccc | ctccactacg | 420 |
| aatttcaact | tttggggagt | gtataaggca | caactcccac | tgaat | | 465 |
| <210> <211> <212> <213> | 19995 324 DNA Glycine max | ς | | | | |
| <223> <400> | unsure at a 19995 | all n locati | ions | | | |
| ttgcatgcaa | gctngctgnc | tacagtttgt | tgatgcacct | atatcactat | gggtagtggc | 60 |
| ggagactgca | gttgtttcgc | ccatgacaat | ggcgaaccat | gcatgcatac | atagttgctc | 120 |
| ctgtcacgct | agacctagcc | tgctacccta | cctttgcgcc | taccctgctg | caacgccaaa | 180 |
| cctatcctgc | tcccctcccc | ctgcgtgcct | acagtgcctg | aggtgtatca | tcattggcac | 240 |
| tggcgatacc | ctccccacgt | tagggctgac | ctatatngcc | agttgcagtg | gcaaacatca | 300 |
| ctggctatct | cgccagcgat | agtg | | | | 324 |
| <210> <211> <212> | 19996 576 DNA | | | | | |

| <213> | Glycine max | |
|-------------------------|--|-----|
| <223> <400> | unsure at all n locations 19996 | |
| agcctcgtcg | g tacntcantt ccgaagacac cgcacggtac gcatccgctc actacatata | 60 |
| annaaannar | naagagggnt tttgatgcct cctagacnga ccnatanata ctaaagccgg | 120 |
| aacaaataga | ccgacaagtc gaacaccaca agcaagggct gttctagtat tgacgacgag | 180 |
| acggagcctg | cgaacacgtc gccagacaga agcggcaacc gctagaagat caccttcagc | 240 |
| agaacgcact | ctatcacacg cagcgaccgg acattcggag cgcacgcaac tagaaagacc | 300 |
| aacgctacac | aggcacgcca aacaacacct ctctttcaat aacaaccacg cgcgcaccag | 360 |
| agcgagcgac | ctacaacccg aagagcacgt ccaaccaccc gaagacttaa cgagcagatg | 420 |
| acaatgagct | gccgcaagct tgacctgatc atccgagatc gatacgaaca atgaccacac | 480 |
| acacccatgg | gagacaatgc agcaaagact acgcatactc gtgacaacag gcacataccg | 540 |
| agaccctgga | cagaggaagt gaacggacac cacacg | 576 |
| <210> <211> <212> <213> | 19997 367 DNA Glycine max | |
| <223> <400> | unsure at all n locations 19997 | |
| gccggaggta | ttaattgata ttcaccanca naatgggcac actgaaggaa tgattggcac | 60 |
| acctgataag | aaatagaatg ataactaaac ttactgaata tacaagcagt ttgagaacaa | 120 |
| caacttacta | ctaaatggag taactggaat gaatgtgaga tgtagaaacc ttcaaagatg | 180 |
| tgcacaccca | tgagactttt gtgacagatt atgcagacct cacacccaat gggtattaca | 240 |
| gaattattcc | caaagataca gacaccaagg catgacaaaa tctattatac tgatcacaat | 300 |
| tccctgtcat | accgatacag tatgaacaat acggacagac accggaacca tttataagct | 360 |
| taaagta | | 367 |
| | 19998 442 DNA Glycine max | |

| <223> <400> | unsure at 19998 | all n locat | ions | | | |
|-------------------------|------------------------------------|--------------|------------|------------|--------------|-----|
| agcttaataa | atttcctatt | gatccacatt | gtgtatgtat | gattgcatgg | aatgagatga | 60 |
| aatgcaaaat | tgggaattga | ttgttagttg | tttggatgaa | caaacactta | . cctgaaacac | 120 |
| ttgtgtgctt | gagataaatg | ttggctttgt | gaggaaagaa | gcttagttaa | ccttcctgga | 180 |
| agcttgacat | acttgctaac | cattttcatc | tctaaagagt | attattgcat | gcttctatct | 240 |
| tgagattatg | acaaatgcta | atttggggga | gatgatgatc | tgtgaaatgt | atgcagtcat | 300 |
| ctcagatatt | gtggttggta | ctttctgaac | aggtcattaa | tctaacttag | catagttagc | 360 |
| tctcttttgc | ttgngacaag | cacaactcta | aatttggggg | agtttgataa | ttgatgtaca | 420 |
| taagtagatt | atgttattaa | aa | | | | 442 |
| <210> <211> <212> <213> | 19999 474 DNA Glycine max | × | | | | |
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| tctccctatg | acaacccacc | attcatcttt | gctggagaaa | aagtcaatat | ctaaaggatg | 60 |
| ·catgtggtta | tacaccaagt | ttggtcatga | ggggtcattg | tagaaggacc | taaaaatcca | 120 |
| atatttgata | gtgcacgccc | acacttncta | caacatacta | cttagtagac | catctntgaa | 180 |
| tgtgatcggn | gctatagtgt | tcacccccat | ttggctatga | nattccccc | aaataaagga | 240 |
| tccatcataa | cagtccacaa | cgaccagagg | aacgttaggg | aaggttacat | ggtaagcctc | 300 |
| aagattaacc | gtttggtaaa | aagtgagtcg | tgtcgaactg | tgcactatgt | agaaattatg | 360 |
| gtagggacct | gttatgactt | ttggcaagcg | taccaattgt | cactatagat | tntacatnta | 420 |
| taaaagagtt | cgtctccaca | gggactcgag | ttacttaatt | cattcggata | aagg | 474 |
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| <400> | 20000 | | | | | |
| agcttgatca | ttacaattat | ctaatcattc | caatccactc | aaatcattca | attgctcatt | 60 |



| cttaagaact | catttacaaa | atggatctan | ggcttatcg | t aggatagcgc | gcttagcgct | 420 |
|--|--|--|--|--|--|--------------------------|
| gctataataa | a | | | | | 431 |
| <210><211><212><213> | 20003 452 DNA Glycine max | | | | | |
| <400> | 20003 | | | | | |
| tcgagaattg | cccaaactcc | ctctccattt. | ctgatttagg | g cttttattgt | ggccttgttg | 60 |
| gtgcttgtgt | gcttagcgca (| actctggctc | gcttagtgcg | r cattagtgaa | tttcggctta | 120 |
| gcgctcgtct | tttcgcttag (| cgaatggact | taaatggtgc | acttaacgag | attagccctt | 180 |
| gctcagcgaa | catgcatage (| tcattcttct | ttcagattct | tcctcgcgct | cagccaaagg | 240 |
| agtgttgcac | tcagtggatg o | gctcgctaag | ccaaaagatt | ggcttagcga | gcggatgata | 300 |
| attagcattt | cacagacttg (| cctaattaac | ctgaaattga | gaggaaatga | ttgttaaaca | 360 |
| cacaaaatgg | gagtactaag t | tatttattac | ctatctttaa | caaaaagtaa | ttacaacact | 420 |
| acaaaataac | cataaattgg a | aggaatttga | ta . | | | 452 |
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| <223> <400> | unsure at al 20004 | .1 n locati | ons | | · | |
| gagggacttt | | | | | | |
| tagagagaaa | gatganctca g | acctccana | accaaaacta | gtcgtagaga | ccaggttatc | 60 |
| | gatganctca g | | | | | 60 120 |
| | | gtcaagagc | ctaagcaaga | ttggactgat | gagacatcgc | |
| ccaaagaccg | agtagttttt a | gtcaagagc | ctaagcaaga gggcctacac | ttggactgat actcttccca | gagacatcgc ctacatacta | 120 |
| ccaaagaccg gtgcactcac | agtagttttt a | gtcaagagc | ctaagcaaga gggcctacac tctatactgg | ttggactgat actcttccca tctcccacct | gagacatcgc ctacatacta cacacactga | 120 180 |
| ccaaagaccg gtgcactcac aaataccctc | agtagttttt a tcattatcct c catctttgac t | gtcaagagc tttttgata gtgaccagg ccatcataa | ctaagcaaga gggcctacac tctatactgg ccgtcctctc | ttggactgat actcttccca tctcccacct agactagaga | gagacatege ctacatacta cacacactga gactgtaggg | 120 180 240 |
| ccaaagaccg gtgcactcac aaataccctc | agtagttttt a tcattatcct c catctttgac t caataatcga t | gtcaagagc de tttttgata de gtgaccagg de ccatcataa de tactagccg de | ctaagcaaga gggcctacac tctatactgg ccgtcctctc | ttggactgat actcttccca tctcccacct agactagaga agagacgcga | gagacatcgc ctacatacta cacacactga gactgtaggg gccatctgtg | 120 180 240 300 |

| gatgagcg | | 488 |
|----------------------------------|--|-----|
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| <223> <400> | unsure at all n locations 20005 | |
| agcttctatt | ttagctgaac cattntatca ataaacacac gttgagtttt attcagacaa | 60 |
| ttagagttta | tetettttat ettagtgaga gtgattetee taaattettg agtgatteaa | 120 |
| gaacaccttg | gctgtatcaa aggactttca caacctttgt gtgttgccct cgctggaaag | 180 |
| agtgattctt | tccttccttt catcatcacc cttgatcttt caaaccacaa ttccagacga | 240 |
| tccacctctg | cccagaatta tctcgaggcc ataactccca ttttacgcac tcaaattaag | 300 |
| tgattcttga | gcctaaattg aatttcagaa cgagaccttt cacctcgttt tgaatcacct | 360 |
| catttggagc | cctgtagctt cagatatcgg catttctata tttctgtcca cgcaccactt | 420 |
| aaccta | | 426 |
| <210> <211> <212> <213> | 20006 449 DNA Glycine max | |
| <223> <400> | unsure at all n locations 20006 | |
| tgccttgccc | catgatatat ntgagggact tatgatcact atgtttgact aattccttgn | 60 |
| gataaaggta | gtgttgccat gttttcaaag cccgtactaa ggcatacaac tcctaatcat | 120 |
| aagttgaata | gttaagggta ggaccactta gcttttcact aaaataagca attggatggc | 180 |
| cttcttgcat | caacacagee ecaateecaa catttgaage ateacaetea atttcaaaag | 240 |
| aattttgaaa | gtttggcaac gcgagtatgg tggcattagt tagcttttgc ttaagaacat | 300 |
| tgaaagcttc | ttcttgtttc tctccccatt tgaaaccaac attnttcttg agcacttcat | 360 |
| tgagaggtgc | tgccaatgtg ctaaaatcct tcacaaatcg tctataaaaa cttgctaagc | 420 |
| catgtgtcgc | aacctaccct tcggcggga | 449 |
| <210> | 20007 | |

| <211> <212> <213> | 440 DNA Glycine ma | x | | | | |
|----------------------------------|------------------------------------|-------------|------------|------------|------------|-----|
| <223> <400> | unsure at 20007 | all n locat | ions | | | |
| agcttccaca | atatccaagc | aatttaatat | ccaaacatca | tgaactaccc | taaaccaaga | 60 |
| aaacagggca | gaggcagaaa | actctgtcca | aaacacattc | caatagcaca | gctttcccta | 120 |
| ctcaaatacc | ccagtaacat | tctcttcgct | tcgattcgtt | aaccattgga | tcgactcgaa | 180 |
| nattttactg | gaggtcccta | gtacataagt | ctacattttg | accgttggga | tcagctagaa | 240 |
| aattttcaga | acccaatatg | tactaccttt | cccataacca | ataatgcata | agcattttct | 300 |
| gcacaagaac | aaaaattctg | ctgcacaatt | caacaaccaa | tttctgcata | atagggcaaa | 360 |
| tnttcgaaat | ccctcttgcc | cttcatccaa | tttgctcana | ttggatccta | caagtcttaa | 420 |
| atcatgtata | tatcatatct | | | | | 440 |
| <210> <211> <212> <213> | 20008 446 DNA Glycine max | ĸ | | | | |
| <400> | 20008 | | | | | |
| taagctcctt | caactgcaca | aggctcttaa | tatttgaaaa | gtattcttgt | ggaacattca | 60 |
| cccgacgaag | acactgacaa | aaacttatct | tcttctttt | ggacaaggta | tggcaagctg | 120 |
| ggggcaagaa | aattttcttc | ccatcagacc | ttggatgcaa | ctatgatcat | atccccatat | 180 |
| cagctagatc | ttgataggta | ttcaagtcat | ccttcgtctt | gccttgaatg | ttaaggagcg | 240 |
| ttccaatcac | actgtcacaa | aaaattttct | ccacattcat | aacatcaata | caatgtctaa | 300 |
| cgtctagatc | agaccagtac | agaagatcaa | agatgatgga | cctcttcttc | catatgcaac | 360 |
| tattactttt | atccttcttt | tgggtctttc | caaatacagt | attcagggtg | ttgaacccgc | 420 |
| tgatatacct | gctcaccagt | caacag | | | | 446 |
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| <400> | 20009 | | | | | |

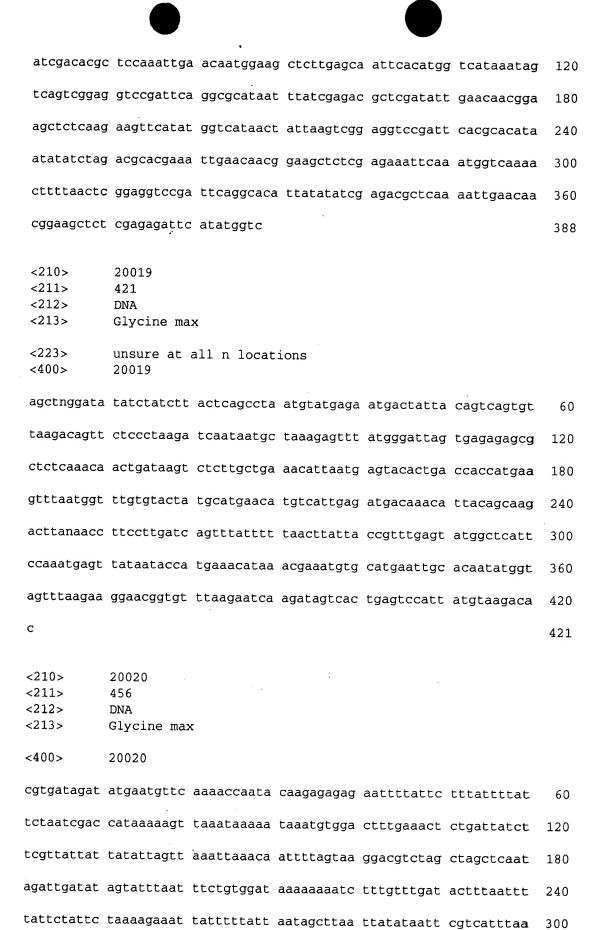
| ttcaagcttt | ggattgttaa | tcgcaccacg | tttcaagaag | agtagagggc | gccacctttg | 60 |
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| ctgagtggtt | atattagcat | tttgttagtt | gaaataaagg | ctcaaacttg | tgttaaagtg | 120 |
| gttgttaatt | ggatttgcac | cacctatagg | cttgttctaa | tttgaagaaa | ttaaggttta | 180 |
| ataaggtgga | aactctaggc | ttgtggctgt | ctcttggctg | accaggagtt | gtgcatattt | 240 |
| acacatgctt | tgtgtcttaa | ttctagtttt | gattaggtat | aatggcacca | ccaattgttg | 300 |
| atattggtga | tcatttcatc | ttctcactat | tgtaaccaac | ttgatgtcat | tcctatttát | 360 |
| aggctacaca | ttttct | | | | | 376 |
| <210> <211> <212> <213> | 20010 427 DNA Glycine max 20010 | ς | | | | |
| tcagacaaca | tgtcataata | gattacgatg | gacctgtaat | taattataac | aaagagtttt | 60 |
| tgcctcttga | agaaactttt | cttcacacta | accatgatga | tgaatgatgc | aatatagata | 120 |
| tcatatgtac | taagatgcaa | catacaagat | aaaaaccaat | acaaatgcca | ctcaagggaa | 180 |
| ttaggcatgt | aaaagtcaaa | acatcttcaa | aacttcttca | agcttttcct | tgaaaggttg | 240 |
| attaccatgt | ggctcatatt | gctccttcta | tctctaacaa | tgtcatcaca | ataaatggca | 300 |
| tggaagtttg | gaaggtaaag | caacatacac | atatgcatcg | tactaccact | cattatgata | 360 |
| tcaatattac | acacttattt | tcctcgtacg | tcactatcac | cttcatagag | tataataatc | 420 |
| atgtcta | | | | | | 427 |
| <210> <211> <212> <213> | 20011 361 DNA Glycine max | | | | | |
| <223> <400> | unsure at a 20011 | ll n locati | ons. | | | |
| ttaagcttgt | aatgttaaga | aaagagcaac | gcacacagtc | atataatacg | gaacaagtat | 60 |
| taaaaaaaac | atataaatat | aáaataacaa | aacaaatcaa | aacaaaaact | ataagcatat | 120 |
| | | | | | | |

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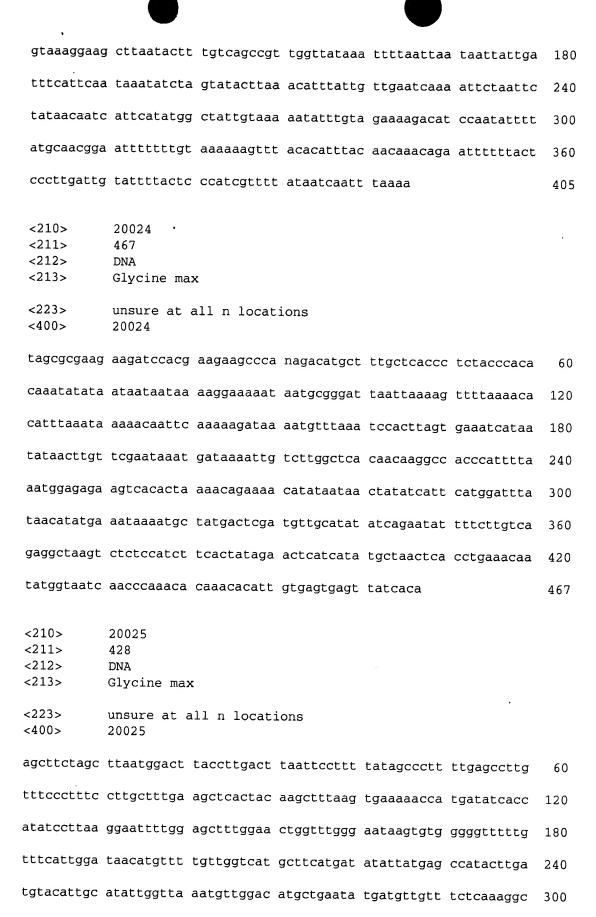
| atagagctg | g aataaaatga | caaaggttga | tctatggatg | g aatgctttct | tagaacctaa | 240 |
|----------------|-------------------|-------------|------------|--------------|------------|-----|
| gcttttgcat | actatagaaa | ccatgaattg | attgcagcco | aggctcgtta | caagcctaaa | 300 |
| aaagtcctt | agattcagtt | ngtgtgttct | cgactatatg | gcaagagatg | aattgaaaag | 360 |
| a | | | | | | 361 |
| <210> | 20012 | | | | | |
| <211> | 402 | | | | | |
| <212> | DNA | | | • | | |
| <213> | Glycine ma | x | | | | |
| <223> | | all n locat | ions | | | |
| <400> | 20012 | | | | | |
| ctgagatggt | atctggtctt | agaanttaat | ttgctaagat | cctattctgt | gcgattggcc | 60 |
| aacctgagga | atggtgtagt | cttgctgctg | actacttgca | ttgtggcccc | ctgcagctcc | 120 |
| cattcatata | cctagggatg | cctataggtg | ttaaccctag | aaggaaggtg | gtgtgggagc | 180 |
| ctataatcag | aaaanttgaa | gccaaattga | acaaatggaa | ccacagaagc | atctctatgg | 240 |
| ctggcagaat | taccttaatc | aatgctgtct | tgacagcttt | gcccttgttt | tatatgtctt | 300 |
| ntttcagggc | cccttcagca | gtcatcaaga | ggctcactac | tatccaaaga | caatttcttt | 360 |
| ggggtggaaa | cttggaagga | aaaaagatag | cttggatctc | at | | 402 |
| | | | | | | |
| <210> | 20013 | | | | | |
| <211> <212> | 444 DNA | | | | | |
| <213> | Glycine max | : | | | | |
| | | | | | | |
| <223> <400> | unsure at a 20013 | ll n locati | ons. | | | |
| (400) | 20015 | | , | | | |
| agcttattat | tgtgacgttg | aatganatgc | attntgtcaa | agttntgtta | aagagctaca | 60 |
| aattanagtg | tggcttanat | tataatatat | tatttttta | taacagagtt | gaaagatttg | 120 |
| accgtgcact | ntatatatcc | atctttatga | gtgaaatttc | ttcaaatctt | anataaatga | 180 |
| aaaaaaaag | agccaatata | agctataaat | taattaagaa | aggaagaaac | ctctcttctt | 240 |
| ggaattggat | gactactgag | ttgtgatttg | tctttaatta | cttgcgacta | tcccatgcca | 300 |
| ataaactgaa | agtgagaaaa | gagcagatta | aggaagaagc | aagctaaaca | tgatngaagg | 360 |
| atatggatat | ggatatggat | ggcgtactac | cattccgtag | aaaaccaaca | tggagtaacg | 420 |

| aagcangata | atactgagag catg | 444 |
|-------------------------------------|---|-----|
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| taataaggat | ttcgattcaa ctatggcaga atcaagatat ccttttccct cctcttccac | 120 |
| attccctaat | tattgctgac aaacttgatg attcaaattt tttgctgtgg tgacagcaaa | 180 |
| ttgaaccggc | gatatcatct cacatcatct tcagtgttgt gttgtaaatc cagcgattat | 240 |
| gtgtggaaac | caagtccagc aatctgtggc atgctagact tggtcatcca aactttcatg | 300 |
| tactgaaact | tgttctacaa cactgtgacg tatgtacctc ctgttaataa aaatgttgat | 360 |
| gtctgtgctt | cttgttgcga tggccagtca catacactcc cttcatctcc tgccactact | 420 |
| acctatggtg | ctccttcgga atcgatttta gtgaccc | 457 |
| <210> <211> <212> <213> <223> <400> | 20015 411 DNA Glycine max unsure at all n locations 20015 | |
| agctttacca | ttaatttata ttgatcagga gtaaaaaggg ggaaatcttg atctcctcag | 60 |
| tctttagtaa | caagcaccgt cttttactgt atcgttatct acttttcaat gagtgtgcac | 120 |
| ttttgtccat | taactaaatt tcatataaga acaactaaga aatgaatagc aaggttactg | 180 |
| agttgaatcc | tgtgatccta caatatcaaa actcatcctg cagtggtagt attgcttaga | 240 |
| atcatcgtag | cgagtatgat gtgattatta ttacctatca tcacattatc aaagttgttg | 300 |
| taatcaattc | taagtctatt atatgaatgt ntatttgatg taatccatta aattatcatg | 360 |
| gtgatatctt | tatggatgag ctattacgcc aaagctatat taccttctga a | 411 |
| <211> | 20016 460 DNA | |

| <213> | Glycine max | |
|----------------------------------|--|-----|
| <223> <400> | unsure at all n locations 20016 | |
| ntgtacttt | taatgacctg gattggtttt ggtagtgaga tntattgttg gtgtggtagt | 60 |
| tgttgtttad | aagggtagag aatttgaagg gtgtagattg tttgcctggc cagtggccac | 120 |
| tattgtagtg | g ttgcacatgg tgtgcagttt ggtttggctt ggtttggttt | 180 |
| atataattgt | gctttttggt tgcagctggg gccctgacag gagctatagc tggtgctcta | 240 |
| gcagctaaag | ccactaagag tggtcttctc cggggagtta cattgggtgc cattgccggc | 300 |
| tctatactct | ctgtggaggt gttggaagct atccgtgcct attggtgtat ggagcaaact | 360 |
| ggctcacgga | gtgcatcatc tatgtgtggt agtcattctc ttcttgattt ccttggtctc | 420 |
| tttgtgagca | caatgatttg tattaactct cataattatg | 460 |
| <210> <211> <212> <213> | 20017 394 DNA Glycine max | |
| <400> | 20017 | |
| agcttctcga | tctattatgc gcctgaatcg gacctccgag ttaaaagtta tgaccattaa | 60 |
| aatttctcaa | gagetteegt tgattaatte egtgegtete gatatattat gtgeetgaat | 120 |
| cggacctctg | agctaaaagt tatgaccata tagaatatct cgagagcttg cgttgttcaa | 180 |
| tttcatgcgt | ctcgatatat tatttgcctg aatcggacct ccgagttaaa agttatgacc | 240 |
| atttgaattt | cttgagagct ctcgttgttc aatttcgagc gtctcgatat attatgttcc | 300 |
| tgaatcgaac | ctccgagtga ctatttatga ccatctgaat agctcatcag cttccattgt | 360 |
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| ttataattaa | aaattccatt c | gagttcgtca | attattaaaa | cattaaaatc | tcttaattgt | 360 |
|----------------------------------|------------------------------------|-------------|------------|------------|------------|-----|
| ttaaaacatt | tccgttatta t | ttttttgtcc | attacagaat | caattatata | attgagtcct | 420 |
| ttattaaatt | aatgaaattg o | cacatgtgat | cacaca | | | 456 |
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| aacatagagc | ggaggcagag c | cactctacac | atgtctcatt | ccaattccac | agctcttcct | 120 |
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| attcttatta | ttacagtcca t | ctcttttat | cttagtgaga | gtgattctcc | taaattcttg | 120 |
| agtgatgaaa | gaacaccctg t | ctgtatcaa | atgactctca | caacctttgt | gtgtggacct | 180 |
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| agctttgagt | tttntcaacc at | ttgaatcaa a | aattctaatg | tatatgtata | tatctatcta | 60 |
| tatttgttta | ccaatgatcg aa | aattttaat 1 | ttgacaacaa | tgattgattt | ggtactatat | 120 |



| | tacagagta | a aacaaaaa | g aatcgaaaa | a gaaaaagaa | c agcagtagag | g atgagtgaat | 360 |
|----|----------------|-------------------|--------------|--------------|--------------|--------------|-----|
| | aagatcttaa | a atgaccaaa | g aatgatgaga | a ctcttggct | c tactctntat | gtctaaattt | 420 |
| | tatcttta | | | | | | 428 |
| | 210 | | | | | | |
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| • | <213> | Glycine ma | ЭX | | | | |
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| (| cttcttggtg | ggttgatgca | a ctctatctcg | r tagaatggta | ı tgatcactat | cagacatatt | 60 |
| (| ctcaatcaat | tcagttgcct | cttcaagggt | tttcaattct | atcttccctc | ctgctgaagc | 120 |
| ć | atctaacaac | tgcttggttt | gtggtctcag | cccatctata | aacatgttca | attgaattgg | 180 |
| C | ctcagagaat | ccatgtgtgg | gagtetttet | taacaaaccc | cgaaacctct | ccaatgcttc | 240 |
| â | actcaaggac | tcatcaggga | actggtgaaa | tgatgaaata | acaactttcc | cttttgcagt | 300 |
| | | | | | | acgtcttcaa | 360 |
| a | tttttgcct | ttgaatgaat | ggagccactt | cttggcttcc | cctgccaaag | aaaatgagaa | 420 |
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| < | 213> | Glycine ma | x | | | | |
| | 223> 400> | unsure at a 20027 | all n locati | ions | | | |
| n | tcatattca | tatatagatc | attgcaactt | gcaaacttca | ttacaaccag | ccaatgatgt | 60 |
| g | aattcttca | gtttgtgtca | tgtacacttc | ctctttcaac | tcaccattaa | ggaaagttgt | 120 |
| ti | ttcacatcc | atttgccata | tctcataatc | atagtatgct | actatggcaa | gtagaatccg | 180 |
| a | atttatttg | agcattgcca | caggagaaaa | tgtttcgtca | taatctattt | tttgctgttg | 240 |
| a | caatatcct | ttatcaacaa | ggcgagcttt | atacgtctcg | acctttccat | ctgctccaat | 300 |
| ct | tttttcttg | taaacccatt | tacaaccaga | ttggtttata | tcctttgaag | cttcaactaa | 360 |
| aç | gtccatact | ttgtagatct | tcattgattc | tatttcaaat | tccatggcat | ttttccattt | 420 |
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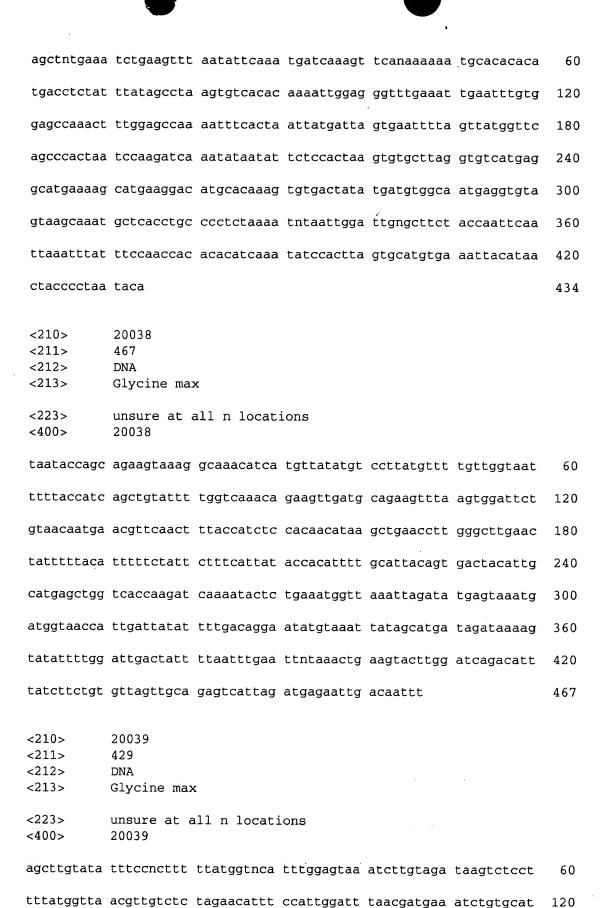
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| ctccataaga | a taatttcagc aaggaaagat gcagtaaaag gagaaatgga gaatcaagtt | 180 |
| agcaaatact | caaacatgtt tgaattttcg attttttatg agtcagatca accggtaatg | 240 |
| | a gaactcatgt gtgtcttaat cacgagctaa tttttcttat atataatatt | 300 |
| gataccaaat | tgacgataaa ttactaattt acttgcaaac gaagaaactg gtatttcaac | 360 |
| ctaaaaggga | caaataattg acgatagcaa tttttataaa atatttaatt atcttaaaaa | 420 |
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| ttagagttta | tetettttat ettagtgaga gtgattetee taaattettg agtgatteaa | 120 |
| gaacaccttg | gctgtatcaa aggactttca caacctttgt gtgttgccct cgctggaaag | 180 |
| agtgattctt | tccttccttt catcatcacc cttgntcttt caaaccacaa ttccagagaa | 240 |
| tccacctctg | cccagaatta tctcgtggcc ataactccca ttttacgcac tcaaattaag | 300 |
| tgattcttga | gcctaaattg aatttcataa cgagaccttt cacctcgtta tggaatcacc | 360 |
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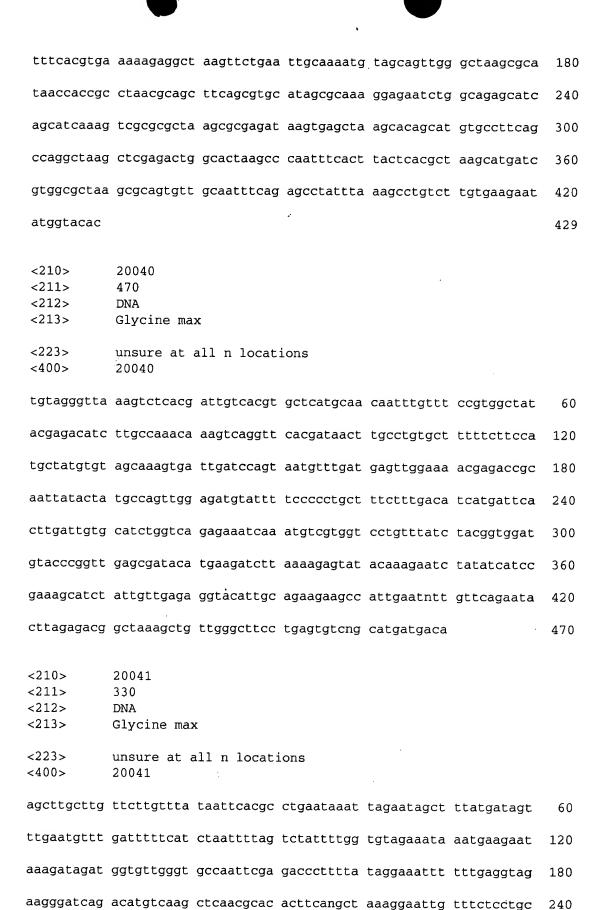
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| aagttgaat | a gttaagggta ggaccactta gcttttcact aaaataagca attggatgg | c 180 |
| cttcttgca | t caacacagee ecaateecaa catttgaage atcacaetea atttcaaaa | g 240 |
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| tgaaagcttd | c.ttcttgtttc tctccccatt tgaaaccaac atttttcttg agcacttca | t 360 |
| tgagaggtgd | c tgccaatgtg ctaaaatcct tcacaaatcg tctataaaaa cttgctaag | c 420 |
| catgtgtcgc | c aacctaccct tctgc | 445 |
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| gcttagtgag | r caaggetege teageeeaat ggetgeegta atgaaatggg ettageeeag | 120 |
| ataggcttga | cttagcgcac gactttcaaa aaaaaattgg actaagttac ccgggcttag | 180 |
| cgattcagcc | tegettagee ecaagtatet caacaggagg atgagtgtte atecteacaa | 240 |
| gatgagcttg | cttagcgcgg taggtgcgct tagcgagttc gtctagaaat gcatatattc | 300 |
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| tcctcattcg | atgatatata actaanaggg actctntcac tctcacttga taggtagata . | 120 |
| gtatattttg | attggaatgt gcggcatcga catgcgttct gcttgatgta atttttaaaa | 180 |
| ttataatatt | ctttctaata tggctactaa tgtacgttta gtggagaata aataactatt | 240 |

| tcttccattn | tccgtccttt | cttcttagat | aaaagacttg | attgctgaaa | ttgagttgct | 300 |
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| tgcacaagag | tttataaatc | gtgagcagca | agttctttct | gtgtacatga | gtgtntttga | 360 |
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| tcttgcataa | gctctggctt | tgttattatt | tattcataca | tatgttggtt | ttgttgacta | 120 |
| tgaatattgt | attatgtgtg | gcaccacttt | ttgtttctct | attaagttnt | aaccttttcc | 180 |
| atatcgttgg | tgattattgg | atatggctgc | aatttaatat | aaaggcctan | agagtgactc | 240 |
| aactatgcat | gtatcgtgta | cattacaaac | ttggggtgcg | tctgaattac | tttatttttg | 300 |
| tatctagaaa | tgctgttagt | caaactacta | gagaacttct | atcgaaaagc | tattttatga | 360 |
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| aatttcgtcc | ggggaccttt | gcttgatgac | atgcgacctt | tctttggtcc | ttgtgaggtg | 120 |
| cttgacaccc | atcattatgc | agtttgcgaa | attccaggac | atgcctaaaa | accaaataaa | 180 |
| tattgatgca | caatccgtaa | gtctacgtga | cacatcggaa | atcaaatgga | agcatcgttg | 240 |
| cataattaag | tgaggttccg | taagtcaaag | aggggatgat | tatgtcattc | tgatggttcc | 300 |
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8390

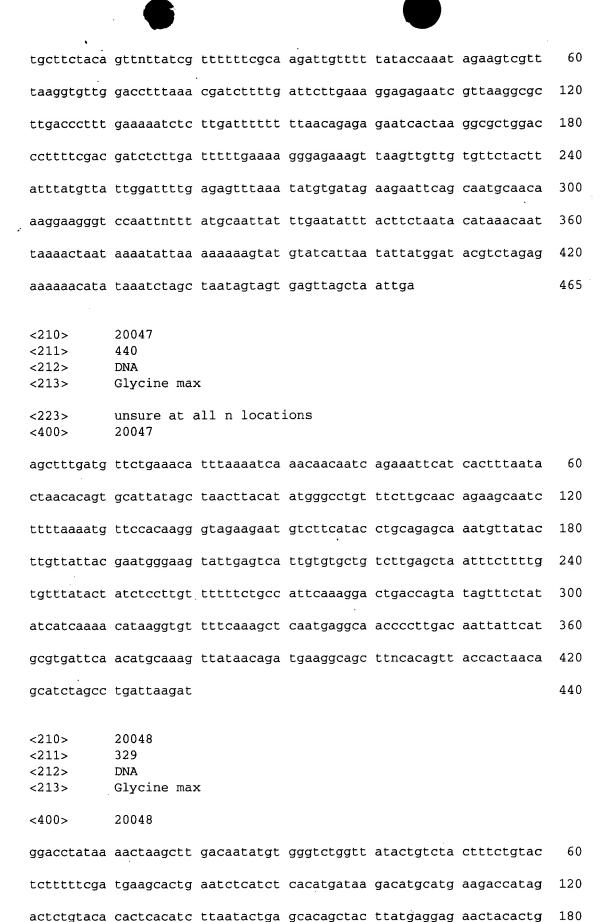
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| ggcttgtggt | ggctggccag ctgtgaatnt tgtgta | aatat gtggattgtg | gcctctggta | 180 |
| atcgattacc | aagggtgggt aatcgattac aaggc | ctata attgaagaca | ggaggctaag | 240 |
| atggtctctg | gtaatcgatt accaaggggt gtaatc | cgatt accaggettg | aaaacgaagt | 300 |
| caggatactt | aaggagcctc tggtaatcga ttacca | agcct gtgtaatcga | ttacacagag | 360 |
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| tctatataag agtttatctc accctggctg attctttcct cctctgccca | ctgaaccatt ntatcaataa acacaa ttttatctta gtgagagtga ttctcc tatcaaagga ctttcacaac ctttgt tcctatcatc tccacccttg ttcttt | gtgt tgccctcgct caaa ccacaattcc | attcaagaac ggaaagagtg agaaaatcca attaagtgat | 120 180 240 |
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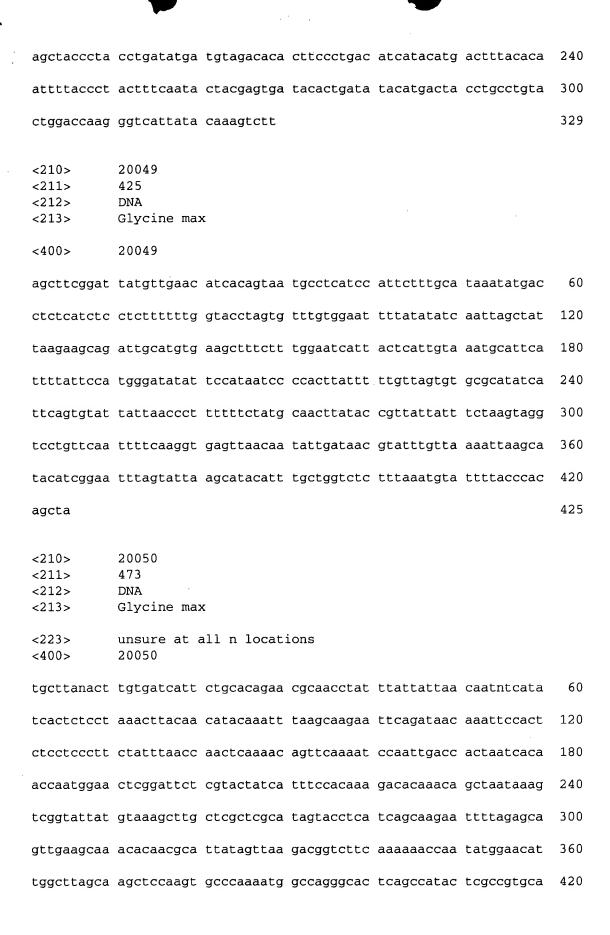


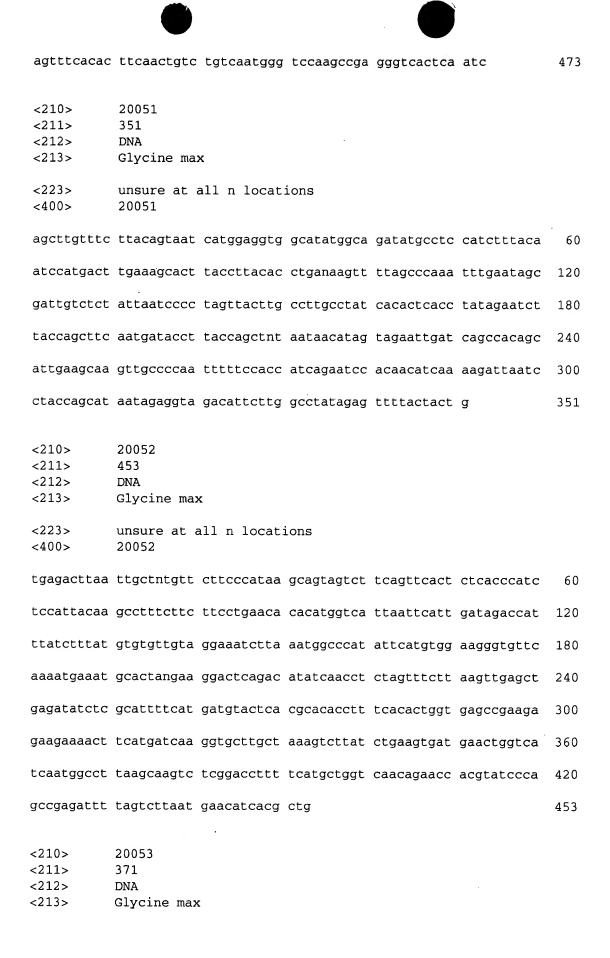


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| gttccgagta | ctttggattt | ggtccgacca | tgccctcctg | atttccagct | gggaaattgg | 120 | |
| cgagtggagg | aacgcccctg | catttacgca | acaagcataa | tgtagacctt | tacggcccta | 180 | |
| aaagctctat | agttgggcct | atgctttaga | gatttcattt | ctggaaggct | ctgtgtcttt | 240 | |
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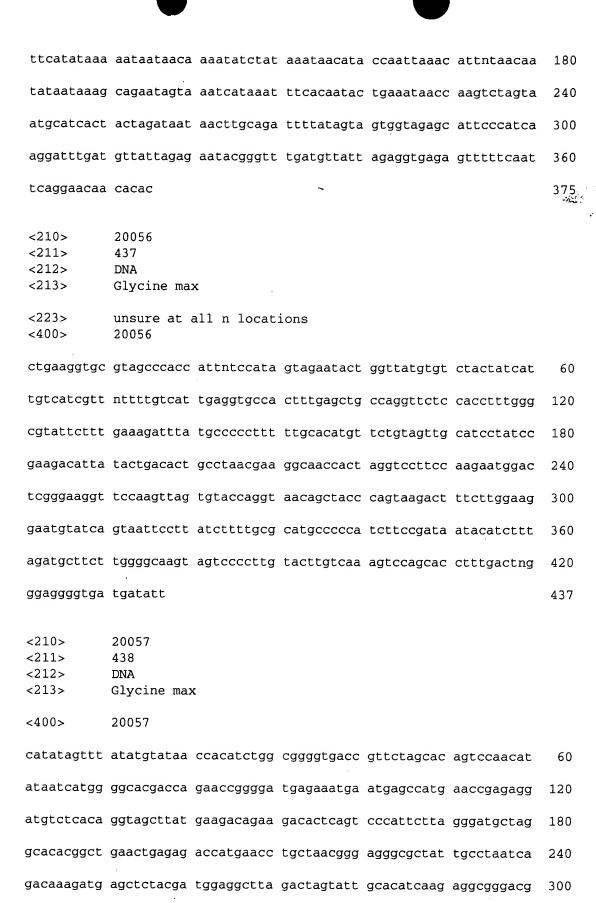
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| aagttcagag | ctgaatagta tttgatgtta tgtttcacca aaatgtgatt tttatagtgc | 180 | | | | |
| tagtgtatgc | ctatattcta tttggatctt atatgagatt tggtgccctt aatattctag | 240 | | | | |
| ttgtatgttg | aaacacatga ttgtaagttt atgttaagca acctgctaga tccccaactt | 300 | | | | |
| tagtttatat | ttcatcattt catgttataa gtatcttatt ctgccattac tgtgttatta | 360 | | | | |
| atccatcatg | cttcgaataa gagagaacct gaca | 394 | | | | |
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| | cccncccc cccccccc nncencnccc ccccccccc ccccncccc | 180 | | | | |
| cccnncccnc | cnecectee ecceencea ecceecece eceneecec etececete | 240 | | | | |
| cccccccc | ctccttnccc cccctccct cccctcccc cctccttccc cctcccccc | 300 | | | | |
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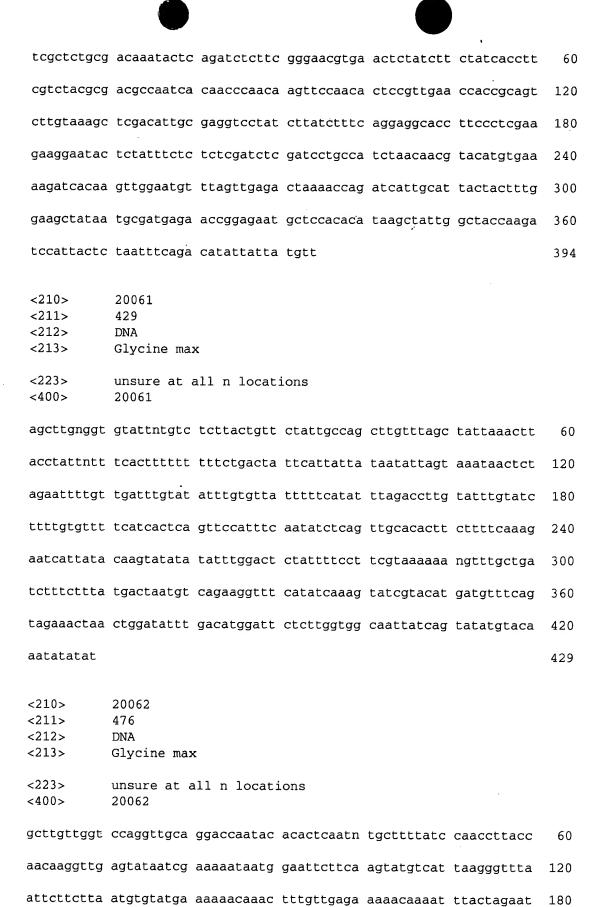


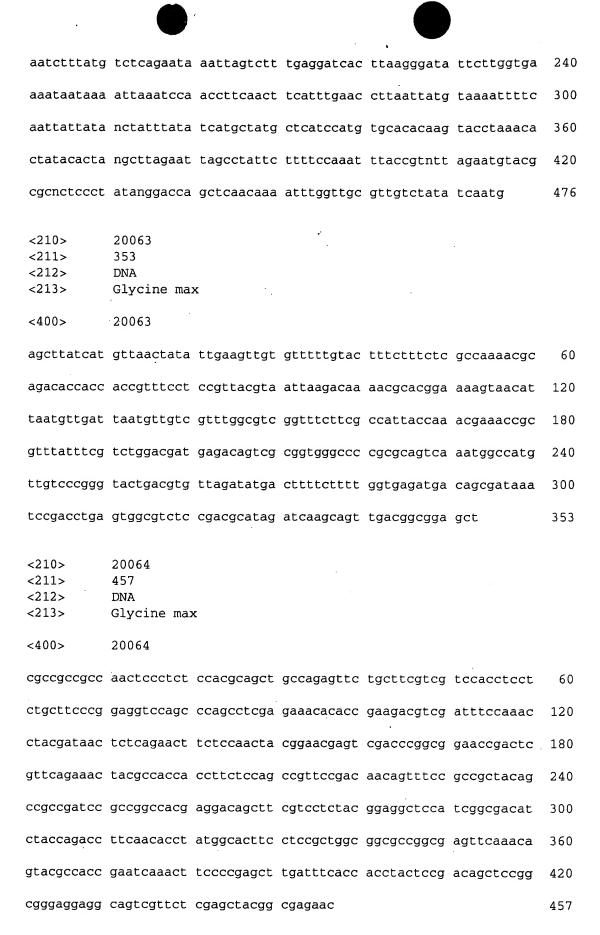


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| gtctcggtta | agcatttaag gacagaggac c | cttaaattct | cttaaggtgt | agacgtggag | 180 |
| cacactgaan | atgaggacac gtagccctct a | aaaggtgagg | gcgtgcagcc | ctctcaagac | 240 |
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| ataactaagc | tcacctcctt aagaagagaa g | gctagagctt | agctacacac | ccctataata | 180 |
| cctaagctca | cttccatgac aaaatacatg a | aaaatacaaa | aaaaaatcct | actacaaaga | 240 |
| ctactcaaaa | tgccctgaaa tacaagacta a | aaccctata | ctgctagaat | ggccaaaata | 300 |
| caaggcctaa | aagaagaata aaaacctatt c | ctaatattta | caaagaagag | tggacccaac | 360 |
| cttgacccat | gggctcaaaa atctacccta a | aggttcatta | gaaccctaag | gccttcttta | 420 |
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| cagtttacgt | agactcgtaa gagttccata g | gactcaactc | gtagacttat | acgagtccac | 120 |



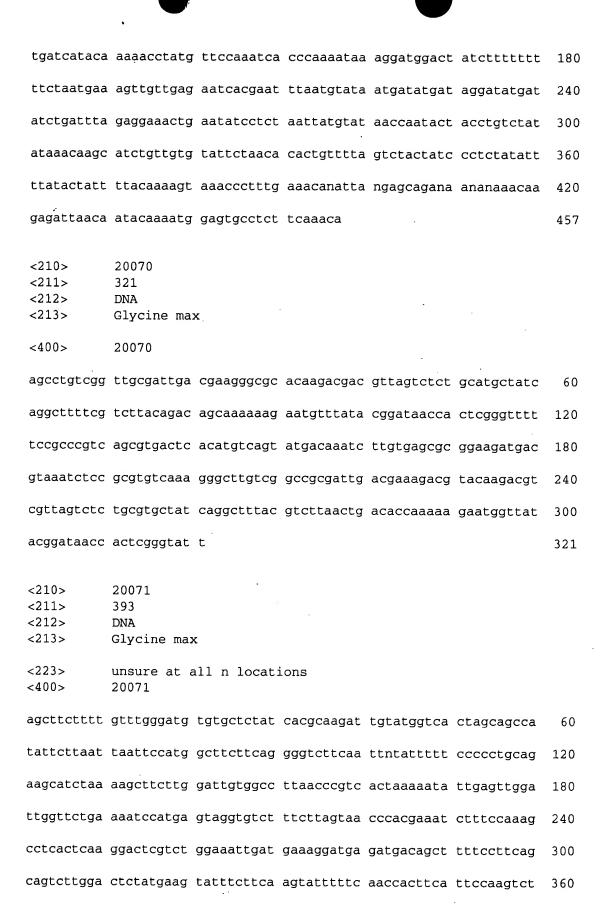
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| gcgatgacga | taattcatat | acaagtgagg | gcctgaatcg | ccatttagct | ggtgacaaca | 420 |
| tacagcacat | atggcaag | | | . *** | | 438 |
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| <223> <400> | unsure at a 20058 | all n locat: | ions | | | |
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| actcacgang | caccaagggc | aacggcgcag | gaggaacgac | agaccaacac | caagagaagg | 120 |
| gtgaaacagc | caacttacct | ggaggacgac | acgacttgag | ggcattgccg | gaagaagaac | 180 |
| ctacgaacca | cgagcccaat | acagctcgtc | gtacatgaat | tctcagagcc | cacacatgca | 240 |
| aaacaacaac | gaga | | | | | 254 |
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| | gttatacgtc | | | | | 60 |
| | | | | | tggccttcac | 120 |
| tcaagaatca | tctttcttcc | actatagaca | actttgtcta | tactatatgt | aacaattcca | 180 |
| gagttaattt | ttggagagat | acctggtgct | tggattgacc | catctataga | gatttgctaa | 240 |
| ttccccttcc | tcattacaag | ctcctttcct | tatctgtatc | tgacttcagg | cacctaaaca | 300 |
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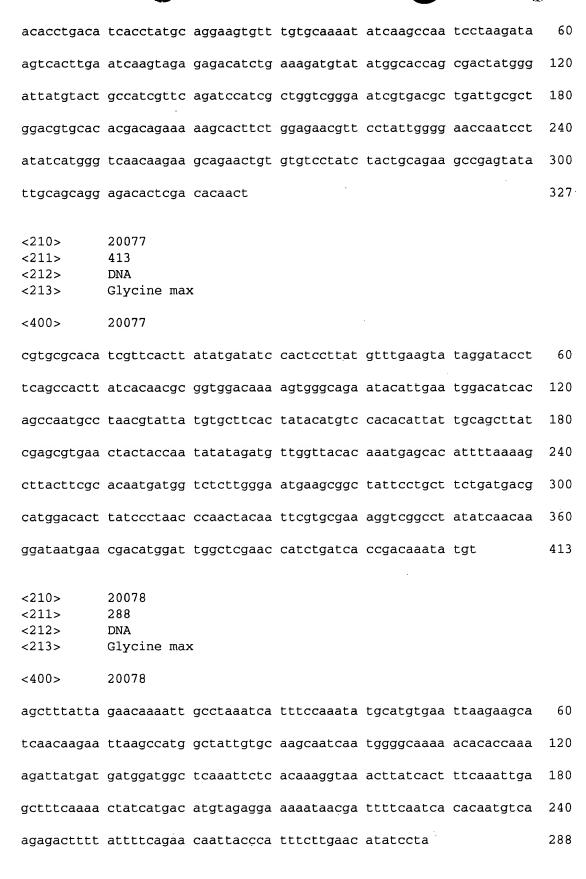
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| tacaaggaca | tccaacacat tcta | gttacc | atacatatat | atatatatat | atatatatat | 180 |
| atatatatat | atatatatat atat | atatat | atatatatat | atatatatat | atatatatat | 240 |
| ntctgaaaag | aacacacatt ctca | tgctca | aggcactgcg | tgaaaattca | cacctaatca | 300 |
| cattctatat | attttgctat caca | cactac | ctacacatat | t-tgaagcaca | tatcataaga | 360 |
| tattcattgt | gtcactcaca ttta | ttata | tgcatatngg | agagctatat | acgtcgtgca | 420 |
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| gtgcttctac | atctcttaat atgg | attatt | gcaaatcttt | gcaagatgag | tttgaaaagt | 180 |
| ttaaaaatga | tcacgatgaa gaaag | gtatga | agttgtaaac | tgagatttcc | tatcttanag | 240 |
| atcatttgaa | taaaggaaag agtga | atctta | gtcacttact | cagtgtgcaa | aagcatacta | 300 |
| ccaataaaac | tggtttgggg tataa | atgagc | aaattgactt | ttataagaaa | actaagttng | 360 |
| caccctccaa | aaaggtgaac ccaa | acaaag | tctccaaaaa | gaanaacata | gt | 412 |
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| ttttgtaaaa | atggagtcat ttttttgt | at tcattccaaa | aaaaacagcc | caacagctgc | 180 |
| aaatagaaaa | caggaggtgc agaaagta | aa ggcccagcag | gtgaagtcag | caataggaga | 240 |
| ggtgacaata | gcaaaagaga agtgggct | ac acgaagccac | gcgcttagcg | cacgtccagg | 3.00 |
| cgctaagcgc | ccaggtacgt tttcaaat | t ttgaatttta | aaattctaag | ggaaaaccaa | 360 |
| gggacgcttc | ccttggtgcg cttagcgg | cc atgtgcgcgc | taagcgcgtg | aatcataaat | 420 |
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| gcaaaacaat | | ıc gtataaccca | ccatcccatg | ttgcccacct | |
| gcaaaacaat | tctccacatc cacaaatc | c gtataaccca | ccatcccatg | ttgcccacct | 120 |
| gcaaaacaat caactgagct catcaatcct | tctccacatc cacaaatc | c gtataaccca | ccatcccatg tcctctcaac acatccaagc | ttgcccacct gccgggtccc atcatgaact | 120 180 |
| gcaaaacaat caactgagct catcaatcct aacacatcca | tetecacate cacaaate cacatactee cacatactee cacatactee cacaacate | c gtataaccca c ttatcctcgt c aagtaattca ag aaaactctgc | ccatcccatg tcctctcaac acatccaagc ccaaaacaca | ttgcccacct gccgggtccc atcatgaact aaccaacatc | 120 180 240 |
| gcaaaacaat caactgagct catcaatcct aacacatcca acaacttttc | tetecacate cacaaate cacatactee cacgtage cecaagette cacaacate agaaaacagg geagagge | ac gtataaccca cc ttatcctcgt cc aagtaattca ag aaaactctgc aa tattctcttc | ccatcccatg tcctctcaac acatccaagc ccaaaacaca gttccaattc | ttgcccacct gccgggtccc atcatgaact aaccaacatc gttaaccgtt | 120 180 240 300 |
| gcaaaacaat caactgagct catcaatcct aacacatcca acaacttttc | tetecacate cacaaate cacatactee cacgtage cecaagette cacaacate agaaaacagg geagagge acacteaatt acceeagt | ac gtataaccca cc ttatcctcgt cc aagtaattca ag aaaactctgc aa tattctcttc | ccatcccatg tcctctcaac acatccaagc ccaaaacaca gttccaattc | ttgcccacct gccgggtccc atcatgaact aaccaacatc gttaaccgtt | 120 180 240 300 360 |
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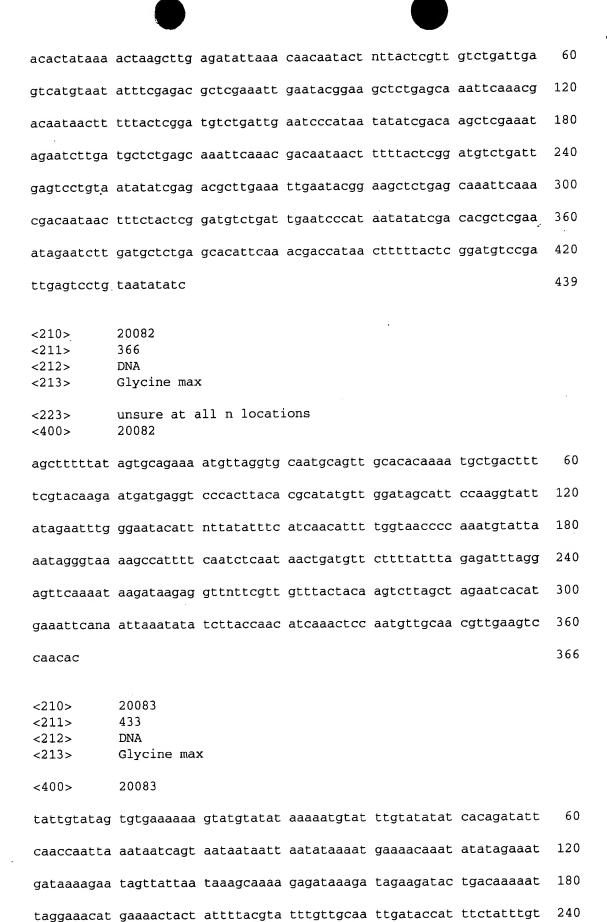
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| atttcttgag | g caaactaatg ctcagacttg gtactttaac taatctttgg tcttc | ttttc 120 |
| tatgacagco | c acaaaatgtt ccaggaatag gtgagttggc tcttggtccg ctcatq | gcttg 180 |
| gggcttcana | a tacactgett aacgetgace attaeggtgt etattteaaa ggaate | gcctc 240 |
| tcttgaatgg | g aaaggtgggc ataccattct cctttatgtg gtaattgggt atctgg | gggta 300 |
| cctgaaaact | tacggttgag tttgtctaat caggtattac ggaggcatgg agatgt | attt 360 |
| gagcgtactg | g ctagaggata ctatcatgca catggtcgtt cggatgatac gatgaa | acctt 420 |
| ggnggaatca | a aggggaaata tttgcctatg tattaattat taaaatag | 468 |
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| tacccttcaa | tcaaagtgct tccattgctc acaggaatga aaagtaaact tgaaag | ggcg 120 |
| cacagagaga | atgacaagat cctacaaaat atggtcaagg atcacaagga aaatga | gaac 180 |
| aagaatgggg | tgacgcacga ggatattatt gatattcttc tcanaactca naagag | agat 240 |
| gacttggaaa | ttcccttgac tcacaacaac gtcaaagcac tcatctgggt tagtat | gcaa 300 |
| ttttctttaa | cattacttta agattcccat gtatacaact atatacgtgc atagat | atga 360 |
| aatttgctga | aataataact tacactttaa tatatataga gagagaggga gagaga | gaaa 420 |
| attgagtagc | gacttattaa taaaata | 447 |
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| gcatgaaata | a cataaccata ctattttagt tatcccaagt atagcaacca agaagtcaac | 120 |
| caattcttt | t gtggtcttta ttaattgtag gtgttaccat acaactaata aagatcacca | 180 |
| ctcaatgtgt | attattttga aagaaagaat ttgatattct gcaaccattt atgcgtaatg | 240 |
| cacccacctt | ggtgcagcaa gtttaaccaa acaaataggc caattcaatt | 300 |
| taagcattca | a tatgacttat gtaattgcat ctaaagcaat gtccaaaacc tcaagtttag | 360 |
| ctcacccttg | g gcatttaatt ntaacaacct aagttgatca ttgtcgcgac ctgccctccg | 420 |
| cgtgggccaa | a gggtgcgtct tccatcanag gaaaacgcgt ggag | 464 |
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| | ggcttagaga atatnggctg tatgatcctg tggaggcttt agtatagaaa | 60 |
| gaataccata | ggcttagaga atatnggctg tatgatcctg tggaggcttt agtatagaaa aagtgcttga tcttagtgga aagcctacag gtgaaggtat tggatgtagc | 60 120 |
| | · | 120 |
| cagagtgggg | aagtgettga tettagtgga aageetacag gtgaaggtat tggatgtage | 120 |
| cagagtgggg | aagtgettga tettagtgga aageetaeag gtgaaggtat tggatgtage tgaaccaata taaateattg gtgttteatt tgetteetta etaettattg | 120 180 |
| cagagtgggg tttgatgttg tttagaagca | a aagtgettga tettagtgga aageetacag gtgaaggtat tggatgtage tgaaccaata taaatcattg gtgttteatt tgetteetta etaettattg egtgttgaag ttttgtttea gaaaaaggtt tttataaaat tggtttaata | 120 180 240 |
| cagagtgggg tttgatgttg tttagaagca gttttgtatc | a aagtgettga tettagtgga aageetacag gtgaaggtat tggatgtage tgaaccaata taaatcattg gtgttteatt tgetteetta etaettattg egtgttgaag ttttgtttea gaaaaaggtt tttataaaat tggtttaata aaacctaett tgttetaaga gaagagaaga tagaaactea tagteaacag | 120 180 240 300 |
| cagagtgggg tttgatgttg tttagaagca gttttgtatc | aagtgcttga tcttagtgga aagcctacag gtgaaggtat tggatgtagc tgaaccaata taaatcattg gtgtttcatt tgcttcctta ctacttattg cgtgttgaag ttttgtttca gaaaaaggtt tttataaaat tggtttaata aaacctactt tgttctaaga gaagagaaga | 120 180 240 300 360 |



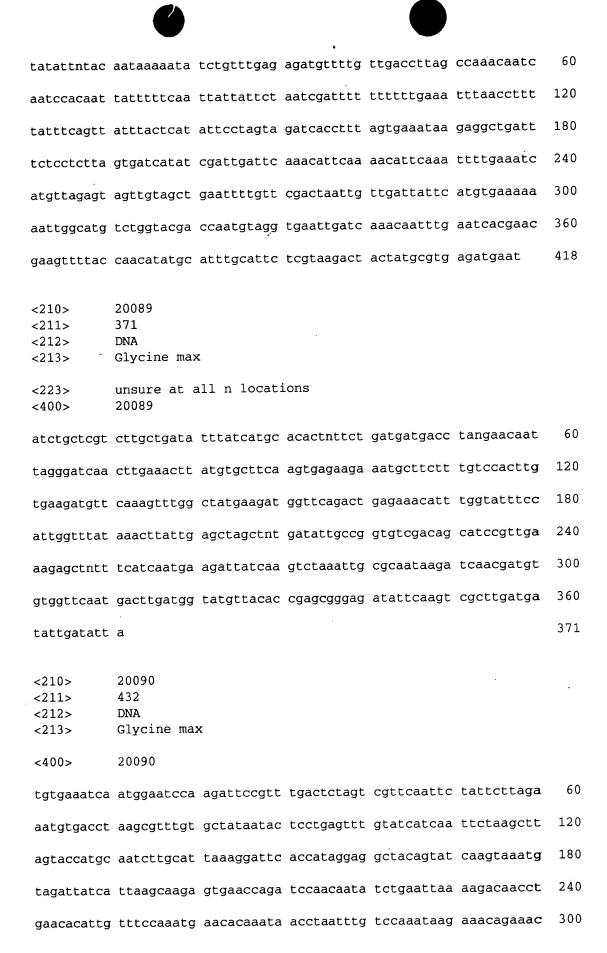
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| ccaaatggcg | gtggtaaacc tagggaagat ctaccagtct cccttccctt | 120 |
| ccccaacgca | acctatgctt atcatggagg tgtctcgggg cattcaatag aacaatgcgt | 180 |
| ggctttcaaa | cacaaggtcc aaagtttgat tgacgcaggg tggctgacat tcaaagagga | 240 |
| caacctaaat | gtgagtacaa aacctcttgc cagtcatggg ggatccgcag ttaatgcggt | 300 |
| ggaggagtat | aaaccttggg ggctgaagca gatggaagat gtggtaacct ctaggaggtt | 360 |
| tatattagaa | tcgttgtgcg aagcgggcat gatttgcctt gacgggcata agggagattc | 420 |
| ttgtttgatg | ca | 432 |
| <210> <211> <212> <213> | 20080 374 DNA Glycine max | |
| <223> <400> | unsure at all n locations 20080 | |
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| gttggatcaa | atggagaata gagatcataa tgaagaagaa aggaggagaa gagggaatga | 120 |
| tggtgttcct | agacaaaacc gaattgatgg tattaaactc aacattcctc catttaaagg | 180 |
| aaagaatgat | ccggaggcct acttggagtg ggagatgaaa atagagcatg tattctcatg | 240 |
| caacaactga | ggaggacaaa aaggtgaagc ttgccgccca cggaatttcc gactatgctc | 300 |
| ttgtgtggtg | gaacaagcta caaaaggaga gagcaagaaa tgaagagcca atggttgata | 360 |
| catggacgga | gatg | 374 |
| <210> <211> <212> <213> | 20081 439 DNA Glycine max unsure at all n locations | |
| <400× | 20001 | |



| ttaactatat | tgttgcataa | tattaatatg | ttttacatqt | acaggacatg | ctctgctgct | 300 |
|-------------------------|------------------------------------|---------------|------------|-------------------------|------------|-----|
| | | aatcttacat | | | | 360 |
| | | | | | | |
| tcaaccctaa | tgtcgacccg | aggtacgtag | ctaacagtat | agtgatactt | gtagataagt | 420 |
| tttcgttcat | ata | | | | | 433 |
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| acttctatag | cctatgtacc | ataggcatca | agctaatcta | t t agagtaga | cgagtagaat | 60 |
| gactccctat | tctgtagtaa | ttatagttaa | ttatgttagt | tgacctcaac | caaccttacg | 120 |
| ttactcgtca | gttactactg | ttagatgtaa | taactgacta | gctgaagaat | aagtactaca | 180 |
| gatgttactg | acttaactag | agtaggatga | tgttacaact | ccattcccga | attctgagat | 240 |
| cacaccctct | tttcattacc | gaaataatgt | tatggcatta | tagagcactt | gattgcatcc | 300 |
| tgagtaacaa | aggctttcgc | aaacacatga | tgggtgataa | gaggagtgct | gctgttggat | 360 |
| actaacaata | ggttgtgaat | atgattagt | | | | 389 |
| <210> <211> <212> <213> | 20085 365 DNA Glycine ma | · x | | | | |
| <223> <400> | unsure at 20085 | all n locat | ions | | | |
| agtttgttgg | attatggggt | acccgtcata | tatggtacta | ggtggcgatc | gggcgatggt | 60 |
| gcaaatcaac | tctcccacat | ccacatatca | aacatgaacc | caccatccct | agttgcccac | 120 |
| cttcaactga | gctcacgtac | tcctacgtag | cccttatcct | cgttcctctc | agcaccgcat | 180 |
| ctctggttcc | agtccctcgc | gtttctctgc | acccgtcggg | gcccgttttc | gatagtaggc | 240 |
| aatatatata | tatatatcan | aacgctcaga | atgaaaccct | ganggtggtt | cagaggttag | 300 |
| gtttgtaaat | ttttagtggc | acgcaaaacg | aataatttta | gactaattaa | ttgagaataa | 360 |
| tctat | | | | | | 365 |

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|-------------------------------------|--|-------------------|------------|------------|------------|-----|
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| gtttgactga | ttacccaagc | cattttgggg | tagggttgaa | tatatggcaa | taattgacac | 120 |
| ctggtgcctt | ccacattttg | aaatatgaca | gaatcgtaac | tataagtgtc | acaattatcc | 180 |
| tgtgtagtac | gttattttga | ttggcataaa | ggaacatgtg | atatgatatt | ttttagtatt | 240 |
| tacatttact | tatcactatg | tatttctttc | aggttaataa | gaaatgtatg | ttttaaattt | 300 |
| ttgcttatat | agtgttaact | gctaaggaag | ctaagctatc | ttatttttat | atagatctaa | 360 |
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| acagt | | | • | | | 425 |
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| catgaaggag | cagatgacaa | cgatgatgga | agcaatgatg | agcatgagga | agatgatgga | 120 |
| ggttaacacc | gctatagttg | gtgttgcaag | cactaatact | gaggtggacc | cgatccaccc | 180 |
| gtccgatttc | aatcgagtgg | gtcgtccggt | ctcggatgta | gtaggccaag | gaggcaaggc | 240 |
| agcagaaaat | gcatggnggc | cccattatgt | tgaagttcag | agcaagcatt | cttttccgcc | 300 |
| atatggtntg | cctcccaatt | ataca | | | | 325 |
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| <223> <400> | unsure at a | all n locat | ions | | | |

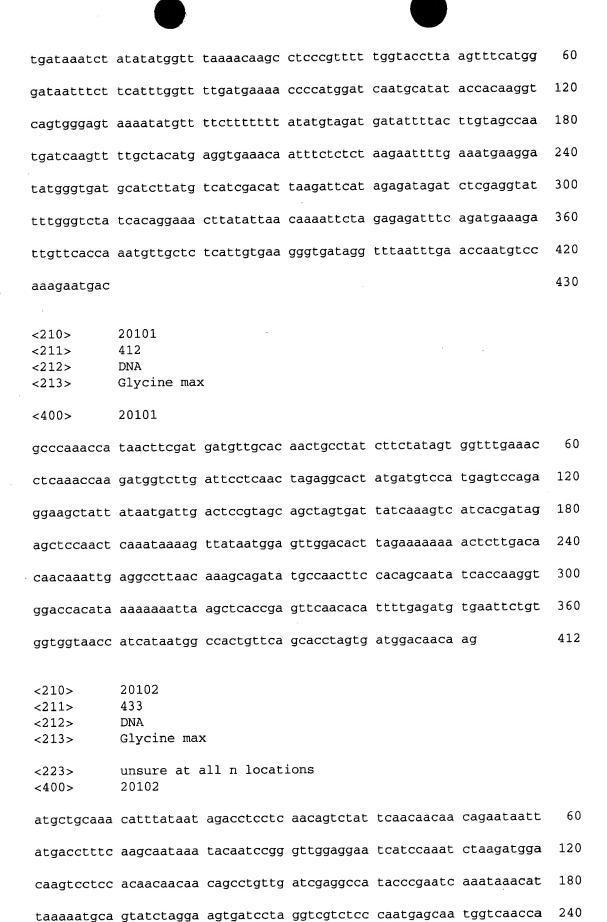


| caaatttcgt | ctaaatgacg | gtacaacaaa | agtgtctttc | acatccaaat | aaaaaacagt | 360 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| acataataaa | aatctaaagt | gccctatagc | tttcacttcc | accgatctac | catctccaac | 420 |
| atagatcaat | ct | | | | | 432 |
| <210> <211> <212> <213> | 20091 409 DNA Glycine max | s. | | | | |
| | unsure at a 20091 | all n locati | ions | | | |
| agcttttcga | ttcattctat | gtacccgtag | tggtccacat | tgtgtttcgt | gcatttttat | 60 |
| tctcgttttg | tttacttttt | ataccccctg | ttgacgtgct | taagccattt | tacttaagtc | 120 |
| gtttctcgct | caacttaaaa | gtaaaataaa | tttccaccga | acgtttgaat | tgtattatcc | 180 |
| attaacttcg | gttaaaataa | attccgaccg | ttcggtcatg | ccgtaaccac | gttggaaatc | 240 |
| aaaaagaggt | aaaaaataat | ataataatca | aaaagacatc | ttttagtaaa | ataaagcgga | 300 |
| aaatcaatcg | gacgttttct | ctttgggatt | tctcattctt | aatcgaattg | attaataact | 360 |
| aaagtgaaac | taaggctaan | atcaactcgc | ctagtcaagc | tcgtccaca | | 409 |
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| tatgcgcata | tttccttaca | aacgttctct | tgcacttgat | attctattaa | ccgaaaaaaa | 60 |
| tgcacccata | tacaatcaag | gcagcttcgt | ttacctagat | tatttacacg | tacttccaag | 120 |
| gtgtatttgt | tacttacatc | acacacctcc | ttggctaaat | tcacacacat | gcatactcaa | 180 |
| agcattttgg | ggtaccaaaa | attgcacatg | tgcacatctt | ggtatttcta | atacctatac | 240 |
| atacacaaac | ttcatgatga | atcttgacta | tctacacaat | aaggtgctac | attttatgct | 300 |
| cttttcaagt | ttttgctacc | taaagccgca | tgcaaattca | agtatatttt | cctttgctga | 360 |
| ctaaaattgt | attcaaatta | aaaggtatac | attctctggt | aatgtatctt | ctttacataa | 420 |
| catgcaacat | at | | | | | 432 |

| | 20093 430 DNA Glycine max | : | | | | |
|-------------------------------------|--|-------------|------------|------------|------------|-----|
| <223> <400> | unsure at a 20093 | ll n locati | ons. | | | |
| agttttataa | tgtagcagtt | ccagggacca | tcgacgaacg | ggcgatcaat | acaaaaagaa | 60 |
| tactcaatcc | atgggaaagg | aatgaaaatc | acacactntg | cctcaactct | gctaaagcaa | 120 |
| ttggatgtac | agtggtcaac | attggcactc | angacttcat | tgaaggaagg | gtatgtttgt | 180 |
| aggccttcta | gtttccaccc | ataaaagcag | aattattgtg | ggcatgtaca | ctgcagaacc | 240 |
| acaaaattta | agatttaatt | taatttataa | atgaaatctg | gtcagatttg | attatttctc | 300 |
| gatcaaagta | attctcaatc | aagttaaccc | ctttttaaa | tgattccgaa | tgctggtaaa | 360 |
| gtatctntat | agcatgctac | attnttttac | agtcaaagcc | tntctctatt | ctttntggca | 420 |
| atgctacaca | | | | | | 430 |
| <210> <211> <212> <213> <223> <400> | 20094 429 DNA Glycine max unsure at a 20094 | | ions | | | |
| tgttgaagtc | taaatagatg | gtgactagac | atttngtgat | tcaaactcca | tagaaggtat | 60 |
| atagcaacta | tttaattaaa | aaacaaccta | agaattcatt | tagttcatat | atttcgagga | 120 |
| gagctactgg | tttgtttgga | gtagtgcatt | cagatgtgtg | tggaccattt | atggttcctt | 180 |
| ctcttggtgg | gaacaatatt | ttgtttcctt | tgtagatgaa | tttagcagaa | tgttgtggat | 240 |
| ctttcttatc | aagtccaagt | caaaaaaatt | ttcaatcttt | aagaatttta | agttacttgt | 300 |
| tgaaaagcaa | tctgaaaaaa | catattaaga | tacttatgac | tgatggtgga | gttgagttga | 360 |
| gtataccttt | aaagagtttg | aagattattg | caaaggatnt | ggcattcaac | atgaagtgat | 420 |
| attaatatg | | | | | | 429 |
| <210><211><211> | 20095 344 DNA | | | | | |

| <213> | Glycine max | | | | | |
|-------------------------|------------------------------------|-------------|------------|------------|------------|-----|
| <223> <400> | unsure at a 20095 | ll n locati | ons | | | |
| tttatctttc | ttgtggggct | tctatggtgg | ctggatcttt | gagcttcaat | gaggtccttt | 60 |
| aatggtgatt | ttccaccatg | gagatgcagc | ggaagacaaa | ggagaagatg | tgagaggagg | 120 |
| caccatccac | tatggaataa | gccatggaag | aaagagcttc | accaccaaga | tgagccttgg | 180 |
| ataagaagct | cggagaggat | gcttcaatgg | aggaaaagaa | agacggagag | aaagagagag | 240 |
| gggggagcac | ganattgaac | gaagatnaag | ggagagaagt | tgaactttga | gttgtgtctc | 300 |
| acaagactct | cattcatcan | agttacaaca | agtggtacac | atgc | | 344 |
| <210> <211> <212> <213> | 20096 432 DNA Glycine max | 5 | | | | |
| <400> | 20096 | , | | | | |
| tctttaagag | tttcatgcag | tacaacctgc | ccactcttca | attatgcaaa | ttgggggatg | 60 |
| caacatctgt | ggtggggccc | atgagttagg | taagtgcata | tcccaacacg | atgcatccaa | 120 |
| agaattcaac | tacatggcta | atccatatca | tcaagggttc | catcaaggag | gacctcttcg | 180 |
| atacaatcag | ggagaaactt | tttcttaagg | ccaaggttgg | agatcccatc | ctgggaataa | 240 |
| tttcaataaa | gatagacaat | ccatccattt | caccttccaa | ccaagggcct | aatctttatg | 300 |
| agaggaccac | caagctagaa | gacactctga | ctcaatttat | gcaagtttcc | ttgtcaaatc | 360 |
| acaagagcac | ggagttagct | atcaagaaat | tggaggtgca | agtgggccaa | ctacctaagc | 420 |
| aactagatga | ag | | | | | 432 |
| <210> <211> <212> <213> | 20097 264 DNA Glycine max | ς. | | | | |
| <400> | 20097 | | | | | |
| tacattgatg | tttgtattta | tgggaggagg | gtgtatgtca | tttttggttt | aaaaagagtg | 60 |
| tcccactggt | aaaactaact | ttccaaatgt | ttgccttctc | acgaaatggc | cccgaggaag | 120 |
| cttgcctcaa | agaggtcccc | gaaagacaaa | gcagccgaag | gaactatttc | cgctccggag | 180 |

| tatgataatc | accgctttat | gagtgctgta | caccaacagc | gcttcgaggc | catcaaggat | 240 |
|----------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| ggtcgtttct | ccgggagcga | cgcg | | | | 264 |
| <210> <211> <212> <213> | 20098 429 DNA Glycine max | · · | | | | |
| <223> <400> | unsure at a 20098 | all n locați | ions | | | |
| actcagcttg | gcaataagta | ctcccgcatt | tatctctgta | tgcattgtat | gtaggtctcg | 60 |
| tcctttgtca | cgggaagccg | gaaggtccat | atcaccttct | taattgtaca | catggggcac | 120 |
| tgcgccccca | aatgcacaag | taagaagaga | taattttccg | ggctctcgtg | tccgtaaaat | 180 |
| gcattcatat | catgcaccac | ataagcatct | cttcataaca | tcataatgga | catatcctgc | 240 |
| atttgtccgt | tatcatattc | cagcctcaca | ttttgcatga | gtcatggcat | catcatgcat | 300 |
| atgcgttcaa | caaacttttt | gatctgcaaa | attgcatacc | atttgttttc | atgtttgctc | 360 |
| atccttgcgt | tntcctctac | aaaacataaa | caaaaaaggg | ggaagcgtga | aacttcacac | 420 |
| tacattctt | | | | | | 429 |
| <210> <211> <212> <213> | 20099 236 DNA Glycine mas | x | | | | |
| <400> | 20099 | | | | | |
| tatctcttca | aatcattttg | aaaaggcacg | aactacctat | atatatgtgt | gtctgatttc | 60 |
| aaaaagcaag | agagagatat | tccaagagaa | cttcattgtc | aaatgctctc | tcaacaactt | 120 |
| ttgggcacac | acttagcaat | ctattaagag | ttcatccaac | aacttcaata | gtaatatcct | 180 |
| tcttttaaag | agagaattct | tettettett | attaaaagag | attgattaac | ggaccg | 236 |
| <210> <211> <212> <213> <400> | 20100 430 DNA Glycine ma | x | | | | |
| ~400 / | 2 O T O O | | | | | |



| aatgttcata | açaaatagta | ataaaatagt | aacgaattgg | gggggggga | ggatntattt | 300 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| taacaatatt | gtttaaattt | tgagttgatt | aatctcatca | cgacatagtt | atctacctta | 360 |
| tttgaacatt | tggttaataa | tcatatatat | atatatatat | atatatatat | atatatatat | 420 |
| atatatatat | atg | | | | | 433 |
| <210> <211> <212> <213> | 20103 376 DNA Glycine max | κ | | | | |
| <223> <400> | unsure at a 20103 | all n locat: | ions | | | |
| agcttttggt | atgatctggg | acctaaccat | ggcagaggtc | tccacagagg | ccattgtctc | 60 |
| cctcgcccaa | tattatgacc | agcccgtgat | gtgcttcacc | tttgtggact | tgcaattatc | 120 |
| acccacagag | gaagacgttg | aagaaatcct | gagatgccct | ctgggaggaa | ggaaaccata | 180 |
| cctattctcg | ggattctatc | cctctttaac | tagaatttcc | aagatatgcc | aaatctcaac | 240 |
| gcacgaatta | agccacagaa | aggaagtcga | aaatggggtg | gttggagtac | caatgaaatg | 300 |
| ctaggaagtt | aaagcaatac | tcttggangt | aaaggcgaat | gggccccgtt | catggacatc | 360 |
| ctcgcacttt | tgatct | | | | | 376 |
| <210> <211> <212> <213> | 20104 433 DNA Glycine ma | x | | | | |
| <223> <400> | unsure at 20104 | all n locat | ions | | | |
| nttcaattca | ttctatgtac | ccgtggtggt | ccacattttg | tttcatgtat | ttttattctc | 60 |
| attttcattt | actttttata | ccccttttg | acgtgcttaa | gccatttatt | taagtcattt | 120 |
| ctcgcttaaa | ctaaaaataa | aataaatttt | ccaccgatcg | tttggattgt | atcatctgtt | 180 |
| aattttggtt | aaaatgaatt | ccgaccgttc | ggtcgtgccg | taaccacgtt | ggaaatcaaa | 240 |
| aaagaggtaa | aataataata | taataataaa | aaaatacctt | ttagtaaagt | aaagcgaaca | 300 |
| atcaatcgga | cgttttctct | ttgggatttc | tcattcttaa | ttgaattgac | taataactaa | 360 |
| agtgaaatga | aggetaaaat | caaactcacc | tagtcaagct | cgtccacaat | aataggtttt | 420 |

| tgaaagtcta | tca | 433 |
|-------------------------------|--|-----|
| <210> <211> <212> <213> | 20105 341 DNA Glycine max | |
| <223> <400> | unsure at all n locations 20105 | , |
| agctttatat | acaccaacag caaatttttt ttcaagcacc acaaactcta actcaaactc | 60 |
| aaaagactca | aacagaagag aagaagagaa tgcatatttg gtagatattc tatgtgcaaa | 120 |
| cagccttatt | taaagggaaa gggaagagga tggtttcccc tattacgttt caactgttct | 180 |
| gaaccttgaa | aatatcagtc atgttcaatt ntttcttcat tatttctgca aatcctatac | 240 |
| ctgaacccta | gcaaagagta caaagccatg catgaacccc ctatactgaa cctaccctca | 300 |
| tgcatgttgc | actgcacggg ttgtatcttg ctagtttcac t | 341 |
| <210> <211> <212> <213> <400> | 20106 408 DNA Glycine max 20106 | |
| | attggttgta tggacaggtt tgttattgta ttgacaaaac accttttttg | 60 |
| | agaggtggat cggttgaagg atgatccatg atttgcatta taggactttt | 120 |
| | tgcttaccct tgtgtattgc atgagcaatg gcaacaaatg aggtatcatt | 180 |
| gttgttgtca | tccctcttga gataactctc aaagtcaaga gaagatcatg aagctcttca | 240 |
| aagtcaatcg | gtttttcatg agtgtgaaga gctgtagaaa cctccttgta ctcagcactt | 300 |
| agcccattta | aggtgtgaat gacaatgtct gcatcgtcca gtggatcatt ggtgatggct | 360 |
| aattcatcag | tcaaagatta tattccatgg agataatcac tcatggat | 408 |
| <210> <211> <212> <213> | 20107 393 DNA Glycine max unsure at all n locations | |
| <400> | 20107 | |

| agettatata | attaaaatct | cacgattttc | acgtgctcat | gcaacaattg | ttagtcgtgg | 60 |
|----------------------------------|------------------------------------|-------------|------------|------------|------------|-----|
| • | | | | | | 120 |
| ctatacaaga | catcttgcca | aacaaagtca | ggttagccat | aactegeetg | tgetttttt | |
| tccatgctat | atgtagcaaa | gtcattgatc | ctgtcaagtt | tgatgagttg | gaaaatgagg | 180 |
| ccgcaattat | actgtgccag | ttggagatgt | ttttcccct | gctttctttg | acatcatgat | 240 |
| tcacttgatt | gtgcatttgg | tctgagaaat | caaatgttgt | gatcctgttt | atctacggtg | 300 |
| gatgtacccg | gttgagcgat | acatanagat | cttangaggg | tatacaaaga | atctatatcg | 360 |
| tccagaagca | tctattgttg | agaggtacat | tac | | | 393 |
| <210> <211> <212> <213> | 20108 434 DNA Glycine max | x | | | | |
| <400> | 20108 | | | | | |
| tcttggatgc | ttactccaga | tacaactaga | ttaggatttt | tgctccaaag | gaggcgaaga | 60 |
| tgacatttat | cactaaagat | accaactttt | gctaaagggc | tatgcccttt | cagcctaaaa | 120 |
| aatgtagacg | ctacatacca | atgactgatg | gaccgagtct | ttaaacaaca | aataagacaa | 180 |
| aacatcaagg | tatatgtgga | caacgttggc | ggtaagtctc | gaagcatagt | ccaacatgtg | 240 |
| gcagatctgc | aagaagtctt | caaggaactt | tacaagtatg | acatgcgcct | caaccctgaa | 300 |
| aaatgtactt | tcggggtagg | cagaggcaag | ttcctcgact | tcatgatcac | tcaccaaggg | 360 |
| attgaagcca | accctaacaa | atgccctacc | atactagaga | tgcacagccc | gaccaacatc | 420 |
| caagaagtct | agaa | | | | | 434 |
| <210> <211> <212> <213> | 20109 386 DNA Glycine ma | x | | | | |
| <223> <400> | unsure at 20109 | all n locat | ions | | | |
| agtttatagt | cttcttgata | gaggaataac | aaagatccta | atccactagt | acctgttgat | 60 |
| aaaatcaaac | aaaacaatgt | aattataatt | ttaattaaaa | aaagacataa | gttttttcaa | 120 |

ctttattttt ttcatttatg tccttttata attgttctct tatttttaaa tttacgtatc 180

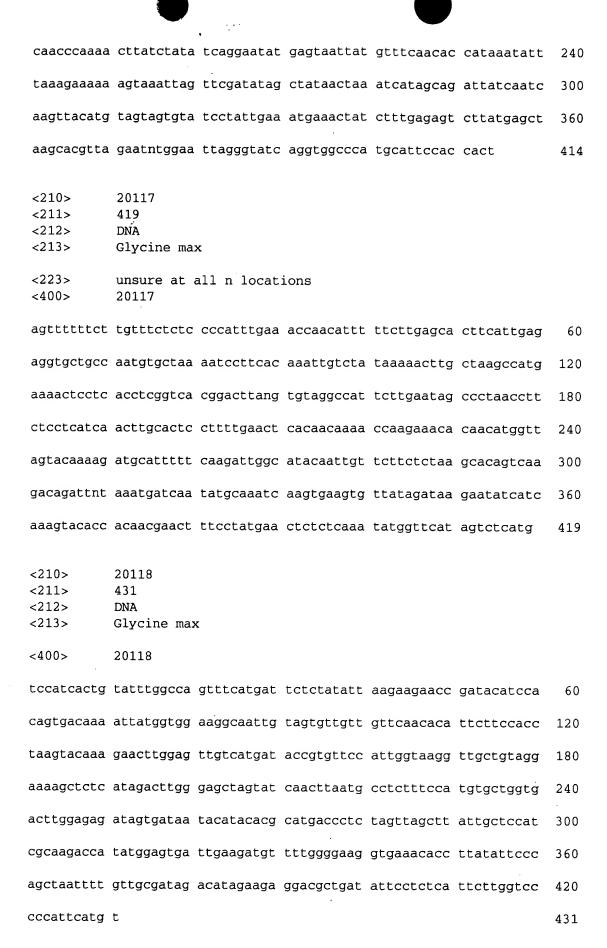
| | | | | | • | |
|-------------------------|------------------------------------|-------------|------------|------------|------------|-----|
| cattaaataa | aaaaaacgta | gtaatcaaga | actgaaaata | gaaaatataa | atttttaaat | 240 |
| aaactaaaaa | taatgtgtag | ataaacggat | aaatcttcaa | tttctaaata | aatggttgaa | 300 |
| aataagaatt | atacctatnt | aatttattag | taaaaatcat | tttaattaac | ttaaaataca | 360 |
| ttntgtcaac | tataaataga | gaatac | | | | 386 |
| <210> <211> <212> <213> | 20110 472 DNA Glycine max | | ions | * | | |
| <400> | 20110 | | | | | |
| aggcgcaatg | atccgtggat | tctcaggacc | ttgaaactaa | gcttggcatt | gtataatttt | 60 |
| ctaaaagatt | attttttatt | ntttatttat | tttttaattt | gtagataaat | aaaattatta | 120 |
| atttttaatt | tatagtagta | tattaaatac | aaaagtagat | ataataaatg | tgaatattac | 180 |
| ttttgagtgt | tcatttttat | attatggata | ttatttacca | tctaattgat | tttttgtgtt | 240 |
| tataattaat | caatttaaat | gtattttcag | taaaattaaa | aaaatttaaa | tttgttaaaa | 300 |
| ttgtgtgatt | tattttatcc | tttcagagtt | atttattta | acaatattgt | ttaaattttg | 360 |
| agttgattaa | tctcatcacg | acatagtttt | ctaccttatt | tgaacatttg | gttaataatt | 420 |
| atatatatat | atatatatat | atatatatat | atatatatat | atatatatat | an | 472 |
| <210> <211> <212> <213> | 20111 377 DNA Glycine ma | x | | | | |
| <223> <400> | unsure at 20111 | all n locat | ions | | | |
| agctttacat | tatattntag | taatgaccca | ctaacctaga | attaaaataa | cttaatgcca | 60 |
| ttaacctagg | gaattaaaaa | aaaaaactta | atggctgagt | gtaactgaaa | ttgtggcaac | 120 |
| caaaagtcac | ccccaacagc | caacaagtca | gccaccattt | ggtctcccaa | aaggttgagg | 180 |
| cctaggttgc | caattgggcc | cttattacaa | cttgaactaa | acctactaaa | gccctttaag | 240 |
| ttgattaacc | caaaacatat | ttttggtcag | ccaactntac | aaggattggg | ccattattta | 300 |
| gacaaactaa | acactctaaa | attgagacaa | agtggtgcca | tttagtcctt | ctccatttgn | 360 |

| gccatgatac | aactcac | 377 |
|-------------------------|--|-----|
| <210> <211> <212> <213> | 20112 425 DNA Glycine max | |
| <400> | 20112 | |
| tgtggattag | gggttggttg tgcatgtttg gcagattttg tagaagcttg tggttaaagg | 60 |
| gtctgttttg | tctttctcat aatctttgaa ggagcttgta gttaaggggt ttgtttttc | 120 |
| tttttcacaa | tatttgaaga agcttgtgtt tgaggtgctt gtttccttta attcagctaa | 180 |
| ccaccttttg | gttgaattcc ctaaaccaat aataagtgtc attttaagta attaacatat | 240 |
| aaaagatgtt | aactaatgta aataaagatt agagacttac caagttactt teettattag | 300 |
| ttgctgcatc | tttgtcattt ttcgtgtgtt gagggataag ttctttctta gcttgattga | 360 |
| ataacatgta | ctatgttgtc attcctagtg actctgctgt caagaactgt gttgttattg | 420 |
| tggcg | | 425 |
| <210> | 20113 | |
| <211> <212> <213> | 359 DNA Glycine max | |
| <223> <400> | unsure at all n locations 20113 | |
| atctttgagc | aaattcaggc gacaatatct ttttactcgt atgtctgatt gagtcccgtc | 60 |
| atataacgag | acgctcgaaa ttgaatgttg aagctctgag ccaattcaaa cgacaataac | 120 |
| tttttactcg | gatgtctgat tgaatcctgt catatatcga gacgctcgaa attgaatgtt | 180 |
| gaacctctga | gcgaattcaa acgacaataa ctttttactc agatgtctga tatagtctcg | 240 |
| taatatatco | agacgctcga aattgaatgt tgaagctctg agcaaattca aacgacaata | 300 |
| actttntact | cggatgtctg attgagtccc gtcatacatc gagacgctca aaattgaat | 359 |
| <210> | | |
| <211> <212> <213> | 20114 427 DNA Glycine max unsure at all n locations | |

| <400> | 20114 | | • | | | |
|-------------------------------------|--|------------|------------|------------|------------|-----|
| tcaacattca | attttgagcg | tctcgtaatt | ttactgtatt | caatcagaca | tccgagtaaa | 60 |
| aatttattgt | cgtttggatt | ggctcagaga | ttcaacattc | aatttcgagc | gtctcgatat | 120 |
| attacgggcc | tcaatcagac | atccgagtaa | aaagttattg | tcgtttgaat | tggttcagag | 180 |
| cttcaacatt | caatttcgag | cgtctcgata | tatgaccgga | ctcaatcaga | catccgagta | 240 |
| aaaagttatt | gtcgtttgaa | ttggctcaaa | gcttcaacat | tcaattttga | gcgtctcgat | 300 |
| atattacggg | actcaatcag | acatccgagt | aaaaagttat | tgtcgtttga | attggctcag | 360 |
| agattcaaca | ttcaatntcg | agcgtctcga | tatattacgg | gactcattca | gacatccgag | 420 |
| taaaaag | | | | | | 427 |
| <210> <211> <212> <213> <223> <400> | 20115 360 DNA Glycine max unsure at a 20115 | | ions | | | |
| | gatctgaaac | tcaacttcct | cttcctccat | ggaagtatgc | tgcgcttgga | 60 |
| | aaataaggct | | | | | 120 |
| aacatcagac | ttgcgatggc | tgtaataaac | ccagtcttca | tcattacaat | ttactctcgc | 180 |
| attggaatga | aaaaatcccc | cagcatttgc | agaagccagt | gttccataac | tgaatgactt | 240 |
| cctaacactg | gaatcctcct | tccccccatc | tgtttcacct | tcctcagaat | catcaagtga | 300 |
| ctcagaatcc | aaaggataat | tgtattcacc | atcctcactc | ctagagcacc | ttncttcact | 360 |
| <210> <211> <212> <213> <223> <400> | 20116 414 DNA Glycine max unsure at a 20116 | | .ons | | | |
| tagccattgc | gaattatatg | cagtcgaaca | tatattatta | tgatctttat | ctttattctt | 60 |

tagtataaac agaaaagatc gactttgatc agtatatgtc ctatggcaat ctattaaaca 120

atttaattaa ttaattattc gacagaatac atatctgcaa gtttcaatat atattttatt 180



| <210> <211> <212> <213> | 20119 404 DNA Glycine max | | | | |
|----------------------------------|------------------------------------|-------------|-------------|----------------|----------|
| <223> <400> | unsure at all n l 20119 | ocations | | | |
| catgttagct | tcaacccact gataat | agga catag | actaa aatac | agngg tagtatt | ttt 60 |
| acgaattcaa | ttgatatatg atagtg | gtca ataaa | catat atgat | ctcta aaagctt. | aca 120 |
| tgttgacatt | gactattcag tttata | ccta tataa | attat ctaat | tctgg tgaggga | attg 180 |
| atgttgaatt | aaaaaaaact aacgga | agat gtaaa | aaatg aaagt | tctct tagccaa | ataa 240 |
| aagaagtaat | ccttaatagc atgtag | aaat gtggg | ttttc tgtct | ccgac cgagtnt | gtt 300 |
| ttcttctaat | tggatcagaa tattgt | aaca aaaat | tgcat tttgt | gcaca ttcattt | ata 360 |
| atatgtaaat | aaataaataa attgag | tctt gtaca | cattt tcag | | 404 |
| <210> <211> <212> <213> | 20120 430 DNA Glycine max | | | | |
| <400> | 20120 | | attta gaag | aagaaa atagga: | aaaa 60 |
| | ctacattgat tgaaca | | | | |
| | agttccttgg tttttg | | | | |
| | cataatatag atttgo | | | | |
| | ctgaaaacag caccaa | | | | |
| - - | ccttgaaacc gtgaca | | | | |
| ggaatcatat | ccaaccaaaa gacaaa | atcaa gtaaa | tgact ccaca | igacaa ccacca | ctgc 360 |
| agaagtgaga | aacggaatgc tttcc | cacca ttggc | tcatc ctagt | aggaa acccgg | cctg 420 |
| agtcaattaa | | | | | 430 |
| <210> <211> <212> <213> | 20121 373 DNA Glycine max | | | | |

| <223> <400> | unsure at all n locations 20121 | |
|-------------------------------------|---|-----|
| tctagctttc | atcaagtggt atcagagcac aagagcttca agtaggtgct ccttaaacct | 60 |
| ccattaattt | tcagctttac cttctcctcc atttttgttt cttcattntt ctccatgtat | 120 |
| ctcctcacat | gacttgtttt gaattttgtt aacatgattt tttagaattt ccaccgatta | 180 |
| aacttgctat | agaagataga tttgattttc tatggttcaa atttcttgct cttgttcttg | 240 |
| aaccacgaat | tgtgttgagt ttaagttcct ttgagtcttg tcttggctat ttttgtgggt | 300 |
| gaaacctaaa | ccatanaatt cttacaaaaa cattaaagta gaagaaaaac ctcaaaaatat | 360 |
| agagtgactt | gtc . | 373 |
| <210> <211> <212> <213> | 20122 426 DNA Glycine max unsure at all n locations | |
| <400> | 20122 | |
| tgaacctcat | cgccactact agatgactcc acttnctatc tttctcctaa agacaacacc | 60 |
| aagttaaaat | caatcatcac aattacttgc catcaaagct actacctttg cccataactt | 120 |
| ttccccttag | gtcataggga gcatatacat taatcacctt tgttggacaa gaggcctcaa | 180 |
| taacttaaga | gggggagaaa ttaagtttca aaatttccca ctaactaact tttaaccctt | 240 |
| ttttaaatga | taggctcgaa atgcagaaga agaagcaaca atcaatttaa taatgttctt | 300 |
| taaacatgca | agacaaaatt gattgcaata acataaatga gataagggaa gagagaaatg | 360 |
| caaactcaat | ttatattggt tcggccactt cacatgtcta tgtccagtcc tcaagcaacc | 420 |
| cacttg | | 426 |
| <210> <211> <212> <213> <223> <400> | 20123 411 DNA Glycine max unsure at all n locations 20123 | |
| | attatggggt acccatcaca tgtggtacta ggtggcggtc gggcgatggt | 60 |
| | tctccacatc cacagatcac acataaaccc accatcccca gttgcccacc | 120 |
| | | |

| | | | | | anagtagtag | 180 |
|-------------------------|---|-------------|------------|------------|------------|-----|
| ttcactgagc | tcacgtactc | ccacgtagcc | cttatcttcg | ttcctctcaa | Cacciggicc | 100 |
| ccatcaatct | ctccaagctt | ccacaacatc | caagcaattc | aacatcccaa | acatcatgaa | 240 |
| ctatcaaaac | caaggaaaac | atggcagagg | cagaaaactc | tgcccaacac | aaacaaatat | 300 |
| cacagctttt | ctcacttaaa | aaccccagta | acattctcct | cgttccaatt | cgttaaccgt | 360 |
| tggatcgact | canannattt | actggaagtc | tctagcacat | aagtctacat | t | 411 |
| <210> <211> <212> <213> | 20124 433 DNA Glycine max 20124 | κ. | | | | |
| | | * | aanaatatat | tagtettta | gatgggatac | 60 |
| | | | | tagtcttttg | | |
| atatgattct | ttcatgcagc | atgatgttca | agaactaaat | cgggttctct | gtgaaaaact | 120 |
| tgaagacaaa | atgaaggtat | ggcaagagat | ttggaatgtt | tgttcatgat | tcttcttgat | 180 |
| gagtgatcat | accaaatggt | tgttgtatgt | tatttttctt | caggaaactg | ttgttgaggg | 240 |
| aactatacaa | aagttatttg | aaggacacca | tatgaattac | attgaatgca | tcaatgtaga | 300 |
| ctacaaatca | actagaaagg | agtcatttta | tggtacttcc | ttatgcattt | tgaattcaat | 360 |
| tatatgttta | gttcttcttt | gttatgtaat | tctaatttag | tttttgcata | tgcatgttag | 420 |
| atcttcagct | tga | | | | . · | 433 |
| <210> <211> <212> <213> | 20125 305 DNA Glycine ma | x | | | | |
| <223> <400> | unsure at 20125 | all n locat | ions | | | |
| agctttctga | cctgagcatt | tgaattgaga | aaaagctcat | tatcttcacg | gcagaattcc | 60 |
| ccaatagcac | gtacatcaac | aaggttccca | gagtctctaa | tttgaagaat | gtgtatagtt | 120 |
| tgatagcgaa | gtgatacaat | tgccaatagg | tcatcataca | agaagacccc | catattatgg | 180 |
| gttanattaa | cgaagtcatt | actgaagacc | ttcttgtcca | agatetetee | atcttccagt | 240 |
| | 225522526 | tanannataa | gaacaataca | taatctagag | tgaacagagt | 300 |

| atgat | | | | | 305 |
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| <210> <211> <212> <213> | 20126 432 DNA Glycine max | | | | |
| <223> <400> | unsure at all n locat 20126 | ions | | | |
| tcaacaccaa | atcctggaca gaatctttgt | : taagtattaa | aggatcaatc | ttattccata | 60 |
| aaccacaact | ctgccaaacc tccaagcta | aagcacaagt | gatgaaagta | ttgcatgcca | 120 |
| acaattcttc | aaggcatctg cacatgaaaa | agttttgtgg | atttatttaa | tcaccatgat | 180 |
| tacaagtitt | tgaagaaaac ttcgatacac | aagccagagg | ataacgcagt | gaaactcatg | 240 |
| taacaataat | tagggggcat ttgttttaag | g atattgagca | agattttaaa | aaatgtgaac | 300 |
| ctaaaaatat | cacattctga tgtttaata | aagttgtaaa | aaataattaa | tttttagaaa | 360 |
| tatattctaa | tgaagaatat tettaaaaa | aaaatttcat | aacttanaac | aagcatcctc | 420 |
| ttaacatatc | at | | | | 432 |
| <210> <211> <212> <213> <223> | 20127 386 DNA Glycine max unsure at all n loca | cions | | | |
| <400> | 20127 gacatacatg aagaaacaa | , gaatttotoa | caaataaact | gtgctcagat | 60 |
| | actttgtccc tttagatga | | | | 120 |
| | ctggtaagaa ttgtgaagg | | | | 180 |
| | agctatagaa ctaacgaag | | | | 240 |
| | ggctcatcca tgttagagc | | | | 300 |
| | gggttatgga agcangagg | | | | 360 |
| | catttctatt cccact | c agaceggaaa | gouoguouug | 909000000 | 386 |
| ayayaayada | | | | | |
| <210> <211> <212> | 20128 429 DNA | | | | , |

| <213> | Glycine max | |
|----------------|--|-----|
| <223> <400> | unsure at all n locations 20128 | |
| tgtgcttcga | ccaacccagc ctcaatttac gtgccattgt gtaagttctt gcttaaaagt | 60 |
| gcaaccttca | atgggaaatg taacaacaag ggtatttcta taacaattga tggcactttg | 120 |
| gtggctcctt | cagattatag ggtcaccgaa aactccggta actggttgga attcgagcgt | 180 |
| gtcaacggag | tttcgattca cggcggggcg cttgacggcc aaggcactgc cttgtgggat | 240 |
| tgcaagaact | ccggcaaagg aaactgcccc agcggagcca cggtatgtta aattaattaa | 300 |
| tttaactctc | atggattgaa cattattcgt tgacatgcac aatttcttgc tcccattatt | 360 |
| attgttntga | tttgatgttg gaggtggagt tgtggacttg gtgattactg agagtttctt | 420 |
| agtggggga | | 429 |
| | | |
| <210> <211> | 20129 390 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <223> <400> | unsure at all n locations 20129 | |
| agcttgtact | ctantgtgtg tgtgtgtg tgtgtgtgat atgtgcagta ngttaacagc | 60 |
| ttctaaaagt | tatgtaaact taggggcaag aggttaacat ttgtaaagga caagcccttc | 120 |
| gtcctctcct | ctccccaatt tcattcacta ctttataagt tggtggaagg acctgctctc | 180 |
| atcacaacag | cagcagccca acagtgtaat taaaatagag acagcttgga aggtgggcag | 240 |
| gggagataaa | tttagatttt gggaggaccg atggttggaa tctgaagcac tgctaatgga | 300 |
| gaaataccca | angctgtacc aaatctcgtg tcaacanaat caaaccataa tgcnagtgat | 360 |
| gagtcacttg | agtagcggat gggaatggag | 390 |
| <210> | 20130 | |
| <211> | 413 | |
| <212> <213> | DNA Glycine max | |
| <223> <400> | unsure at all n locations 20130 | |
| ntgcaacaaa | tagtggctaa tttgagggcg atccaattgt atgaggggac ccaacatcta | 60 |

| gaggcctgca | tgcgtgggtt | gctagttaca | tgtattgctt | gtcgctggag | cagacccatc | 120 |
|----------------------------------|------------------------------------|-------------------|------------|------------|------------|-----|
| aactgcccta | actcttttag | actggtgatc | cctaggctct | tgaccttgac | ttgatagaac | 180 |
| ctctttttaa | gcgaaggtgt | ttgacttgat | cccatgtttt | actaaagtgt | gcgaatcaaa | 240 |
| acttcaacat | ctatcatggg | tgggatggat | gaatgcatga | agaaatgcat | atgacacaaa | 300 |
| tgcaatttat | gaatacggga | gcccgggaaa | ttgtctcctt | cttagataca | acgtcttgng | 360 |
| gtagcacagt | gcccgacgta | tgtatttaag | aaggtgacac | aaaccctcca | ttg | 413 |
| <210> <211> <212> <213> | 20131 344 DNA Glycine max | x all n locat: | ions | | | · |
| <400> | 20131 | | | | | |
| agcttttttg | ttatagtcac | tatttacgta | tcaactatga | tccatatatt | ttcagcatta | 60 |
| tctattatat | tctcttaaaa | aaatcccaaa | taacaataac | tacacattta | ttcatattaa | 120 |
| aagtgaattg | acgtgtttgt | agaattaatt | ttgaattaac | atggctctat | agaattgatt | 180 |
| gtaagtaaac | ataagtttgt | agtatataat | atgattattg | tttggaaatt | tgctaccaaa | 240 |
| actgatattg | tgggcaatta | ctacaatgac | ttgagttcaa | gagtacacat | tgaacaaata | 300 |
| gcatgactaa | taatacgact | caagttagtg | tngattcaaa | agct | | 344 |
| <210> <211> <212> <213> | 20132 412 DNA Glycine ma | x | | | | |
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| | | | | | atgacatcgg | 120 |
| | | | | | aagttttccc | 180 |
| | | | | | gaggatgaac | |
| | | | | | agtttgcaag | 240 |
| | | | | | atgaagttac | 300 |
| aatgatcaat | gcatgtgatt | gcgattgcag | attctacgtc | atcacgtgag | cctatatatg | 360 |

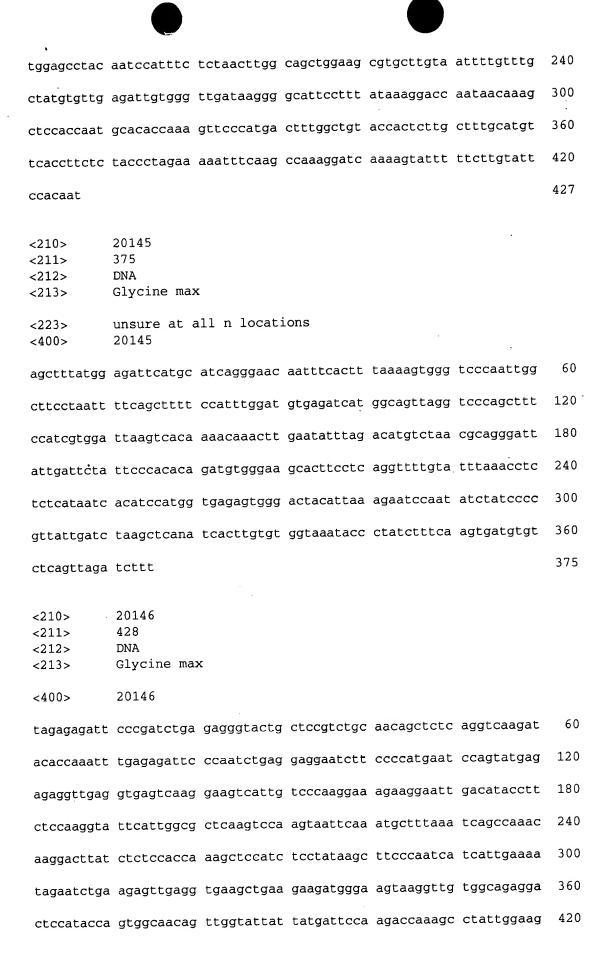
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| <223> <400> | unsure at a | all n locati | ions | | | |
| attctttgaa | gataccgaga | aagatagagc | tgtttgtaat | tttagtcctt | tgttctgttt | 60 |
| gatactccat | aataaaatgg | tatattacta | ttattattat | ttgttctaat | aaatcattta | 120 |
| ctattattat | tttattttat | ggcttgcgaa | ataaaaagaa | gataggaggt | ttttctagag | 180 |
| gtgaatgatg | agaccattcc | atgcctcttc | aattaagtgt | ttttcattga | atctctatat | 240 |
| ttttgtcggg | tcaataacat | atttttgtta | tcagctggtg | atcttattga | tgatgagact | 300 |
| tggcctncaa | ttgtcctaat | cattcatcat | gatattgcca | ataagatacc | gattcatgct | 360 |
| С | | | | | | 361 |
| <210> <211> <212> <213> | 20134 425 DNA Glycine ma | x | | | | |
| <400> | 20134 | | | | | |
| tttcgattca | ttctatgtac | ccgtagtggt | ccacattgtg | tttcgtgcat | ttttattctc | 60 |
| gttttgttta | ctttttatac | cccctctta | acgtgcttaa | gccattttac | ttaagtcatt | 120 |
| tctcgcttaa | cttaaaaata | aaataaattt | ccaccgaacg | tttgaattgt | attatccgtt | 180 |
| aacttcggtt | aaaataaatt | ccgaccgttc | ggtcgtaccg | taaccacgtt | ggaaatcaaa | 240 |
| aaaggaggta | aaaaataata | taataataaa | aaaaacatct | tttagcaaaa | taaagcggaa | 300 |
| aatcaatcgg | acgttttctc | tttgggattt | ctcattctta | atcgaattga | ttaataacga | 360 |
| aagtgaaact | agggctaata | tcaactcgcc | tagtcaagct | cgtccacaaa | aataggcttt | 420 |
| tgaag | | | | | | 425 |
| <210> <211> <212> <213> | 20135 338 DNA Glycine ma | ж | | | | |

| <400> | 20135 | | | • | | |
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| aggccacttg | tccaacaagg | acactatctt | aatgagctta | agcattatct | cgatggtagg | 120 |
| tatatttctc | cttgcgaaag | agaccacttg | ttgttgaaag | attgaacttt | cattttctgg | 180 |
| tttgattcaa | taatatttga | agatgatgac | aacattgatg | ctctgctctc | caagccaatt | 240 |
| gttaaggagt | ccatgtttac | ttggctacaa | gctaatagca | tgttcaatga | aggacaacat | 300 |
| ctaacatatg | tgcaattcat | aacaaagttt | acatatgt | | | 338 |
| <210> <211> <212> <213> | 20136 422 DNA Glycine max | | | , | | |
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| cgttacagaa | cttaggaaaa | atcagaacaa | gcttgttgta | catcgttcgc | gtgtacgata | 60 |
| tccactcgac | aaggtttgaa | gtagaggaga | ccttcaatcc | tataacgcaa | cgtggcggac | 120 |
| aaaaatgggc | agttaacttg | aatggccatt | attgtcaatg | cggaaggtat | tctgcgcttc | 180 |
| actatccatg | ttcacacatt | attgcaactt | gtggttacgt | gagcatgaac | tactaccaat | 240 |
| atatagatgt | tgtttacacc | aatgagcaca | tgttanaagc | atactccgca | cagtggtggc | 300 |
| ctcttgcgaa | tgaaacggca | attcctcctt | ctgatgaggc | atggacacta | attcctgacc | 360 |
| caactacaat | tcgtgcgata | ggtcggtcat | aatcaacatg | gataacgaat | gagatggatt | 420 |
| ga | | | | | | 422 |
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| atcttttctc | taagcttctt | atccaaggca | ctctcttggt | ggtgaaactt | ctccttccat | - 60 |
| ggcttattct | ttagtagatg | gctcctcctt | tcacctcttc | tcctttatct | tctgttacat | 120 |
| ctccatgatg | aaaaatcacc | attgaagggc | ctctttgaag | ctcaaagatc | cagcctccat | 180 |

| agaaggttct | caagaaagct | tccatcagta | aatgtagcca | ttaacttatg | cgtaattnta | 240 |
|-------------------------|-----------------------------------|-------------------|------------|------------|------------|-----|
| atattttaaa | acaataatat | gttatttctt | ccaacaggtt | aagagcatca | ttagtgggag | 300 |
| tatgtactca | atagcaatgg | gagataaaca | tactaaaaga | accccacact | tgtgtatcaa | 360 |
| caatgttatc | acaatatcac | ataaagc | | | | 387 |
| | 20138 414 DNA Glycine max | x all n locat: | ions | | | |
| <400> | 20138 | | | | | |
| tgtaggatta | tggggtaccc | atcacatgtg | gtactatgtg | gcggtcgggc | gatggtgcac | 60 |
| aacaagtttt | ccacatccac | aaatcgcgca | taaatccacc | atcccctgtt | gcccacctcc | 120 |
| aactgagctc | acgtactccc | acgtagccca | tatcctcgtt | tttctcaacc | ccgtgtcccc | 180 |
| atcaatcctt | ccaagcttcc | ccaacatcca | ggtaattcat | catccaaatc | atcacaaact | 240 |
| aaaaaatcaa | gcaaaatagg | gcaaaggcag | aaaactctgc | cccaaactca | aaccaaaatc | 300 |
| acagcttttt | ctcacttaaa | gaccccagta | acatttcctt | cgttccaatt | tgttaaccgg | 360 |
| tggattggac | tcgaaaatnt | actggaagtc | tctagtacat | aagtctacat | tttg | 414 |
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| <223> <400> | unsure at 20139 | all n locat | ions | | | |
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| gcttaattcg | gaactatgat | ggataatgta | attccctggg | agttgatgtc | ttgatggttt | 120 |
| atactttacg | acacttcagt | cgatgacact | tcacttaact | ttcttgttag | ttaaaatctt | 180 |
| tcactaatca | gaatccttcg | taagtagtcc | ttttcgatat | attcataacc | aactatgata | 240 |
| ccactcttag | cacccgactg | ctttcaggat | taatgactgt | ccttgtaaac | catcaaaaga | 300 |
| ctcttttaac | atactttntt | ctcattcaca | cactttccag | agnactctnt | atgagattac | 360 |
| ccatctcata | aatatntcaa | gacaaacaca | cttaactgtg | aaattctta | | 409 |

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|----------------------------------|--|-----|
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| aaattaaaca | aataaaaaaa ccacaattaa aagagggtat gaataagata tgcatatttt | 120 |
| tcaatggatc | ctaacaatgc acaaaaatga gagagataga aaatggaaaa atgatctcaa | 180 |
| ggtaacaaac | aattcatgag cacttgatca caaaatgcaa agtggataag tagttaagtc | 240 |
| aaaattaaga | atgctccatg gtgatagaaa tttttcaaac aacactcaac cctgtcaaat | 300 |
| gtgtatgggt | ttgtgtttac acttaaacac cacatgtaat gatgtaagct ccatttgagc | 360 |
| ttgtatgcct | acgatettet teateaatgg atteattgae ttettgaaag gtgaatgtge | 420 |
| gcagaatgga | | 430 |
| 210 | 20141 | |
| <210> <211> | 386 | |
| <212> | DNA | |
| <213> | Glycine max | |
| <223> <400> | unsure at all n locations 20141 | |
| agcttctctg | agtattcctt aaagcttatt attgacaaaa ctttgcacat aagtttaata | 60 |
| ttgaactatt | tttcaaatca ctagacgaga gtccataggg cccttaatca cacatagaga | 120 |
| tagattacct | aagtggtaat tcgcctatga aggcccaagg tccatggccc aaaaaggttt | 180 |
| gtataaaaaa | gttgagaccc ctgggtaaag accatttete attettgeat ttactateet | 240 |
| atttattgct | tagagtcaaa acttgacttg ggcatcaaag taccttttgt tggtaccccc | 300 |
| ctttggacca | aacactanag caagacgaat agtgtaatga cccgcctcgt cgctacgata | 360 |
| tcacttatta | taaaatgtga catcat | 386 |
| | | |
| <210> | 20142 | |
| <211> <212> | 420 DNA | |
| <212> <213> | Glycine max | |
| | | |

| | • | |
|-------------------------|--|-----|
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| ntaccggtat | ccccattttc agaaaatgga ggtgtttatt caagtcactg ctatgctcgc | 60 |
| tgttggctta | atccaaccta gccatagtct gttctcttct ccgatactcc tagttaagaa | 120 |
| gaaagacgga | acctggcatt aaagggcact aaaggaaatc acggttaaag attgttttcc | 180 |
| tatgccaacc | attgatgaat tactcgacga tttgggccaa gcatcatggt ttttgaagct | 240 |
| cgatttatgc | caaggatttc atcagatacg tatggtcaag acttacattc ataagacagc | 300 |
| ttttcaaatg | cactagggac attatgagtt taaggtcatg ccctttggcc tttataatgc | 360 |
| cccttctact | tctcaagcaa ccatgaatga tgcgctccaa ccatttctga ggaaatatgt | 420 |
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| <223> <400> | unsure at all n locations 20143 | |
| tctagcttgt | tcgaagatcc tcgagacgtc atacgagggg ccaatctttc ggaaaagact | 60 |
| ttcaagaagt | ttttgaagat ttctcttgat gaaaactata acctgcattc ttttgagttc | 120 |
| aaccattccc | acttctgcac catggggttt gttacctggt gggagaaata ttattcgacc | 180 |
| cgttcagttg | gagacactac tatcatgatc tgcagactcg agagtggttn tatacaacca | 240 |
| acggtcgaga | atatccgctc aaaccttcaa gctcgaggta ttaaattcct tttgactttc | 300 |
| tanattgata | tgtactattg gcttctctaa tattcttatt tcaagcaaaa caatcatgac | 360 |
| gaagaanagt | ggctgaacgt ctcgagct | 388 |
| <210> <211> <212> <213> | 20144 427 DNA Glycine max | |
| <400> | 20144 | |
| | g aatccacgaa gcattatttc tcttgtttac caaagttaat gtcgctgatt | 60 |
| | c aaaaagtgac aaaatagctg tactagagta ttgacatggg tattttttgc | 120 |
| taatctttgo | ttgtataatg aaccatgaag accacaggag gcaacctaga gtcagaagta | 180 |

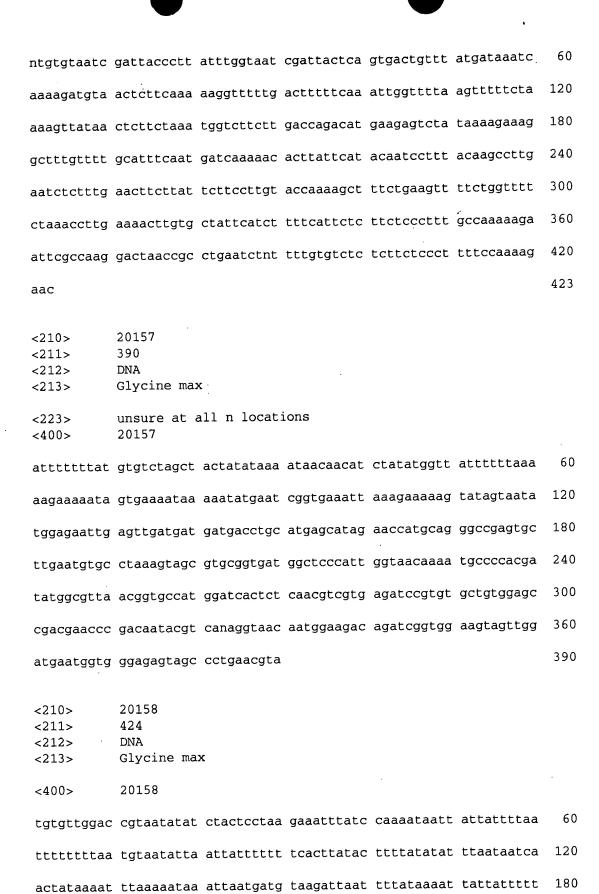


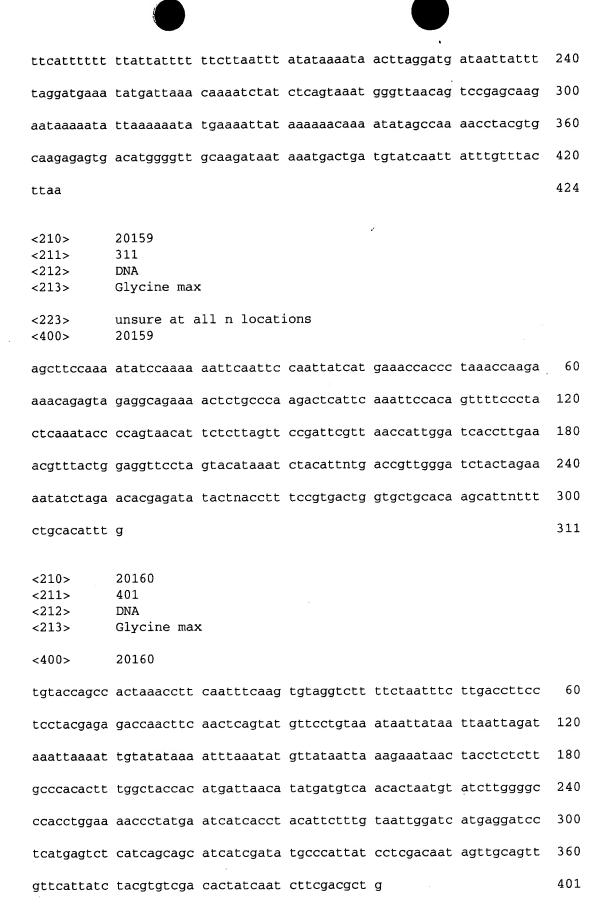
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|-------------------------|------------------------------------|-------------|------------|------------|------------|-----|
| | 20147 375 DNA Glycine max | i. | | | | |
| <223> <400> | unsure at a 20147 | ll n locati | ons. | | | |
| agcttgtctg | gttcttccat | taattggagt | cccaccatt | gtttctgctc | caactctttg | 60 |
| ctctttgttg | tctaggaaga | gcacatcacc | ctcaaaattt | tgaatcctca | acatgataaa | 120 |
| aagactatat | ttttaagcat | ggtggtgcaa | tgaaactgaa | gaacaagagc | aacccatgat | 180 |
| ttanaatgag | ctggagattg | gaaaaatgga | tatagaaatg | agaaaaaaac | aaaatggtac | 240 |
| cttcaaattt | gaaggaagtg | gtggtgtgga | tgtgagtgat | aatggtggcc | agttctggtt | 300 |
| tttggtgacc | attttgaagc | acaagcactt | atgtagagaa | gcatatgaga | gagagagaga | 360 |
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| cctacaattg | atatggtcat | acatccatca | aatatctcta | attaaaatga | aattatgcgg | 120 |
| ttcattacag | aaaataaaat | atttactttt | tatatgagaa | aatcaaatgc | tgatattatg | 180 |
| aatacatata | taataaacta | gagtaacacc | cgtgctatgt | agaagtaggg | gaagaaaaaa | 240 |
| gtttcaaata | gttgctccat | acttttaatt | aaagagagtg | tagagtaata | aaaaaaaaat | 300 |
| tggtgtgtag | gccaatacga | tttttattt | tttacagata | tcaaccataa | ataaatgtat | 360 |
| gtaattaatt | aagctagcaa | catttgtctg | gttcaaagta | gtcctgaaca | ttaaattctt | 420 |
| ctacaattat | tggtgtacat | ttat | | | | 444 |
| <210> <211> <212> | 20149 292 DNA | | | | | |

| <213> | Glycine max | | | | | |
|----------------------------------|---|-------------|------------|------------|------------|-----------|
| <400> | 20149 | | | | | |
| tgagtatttt | tcccttcact | tttttgcttt | ccattttata | aatttgtcat | attcttgata | 60 |
| aaatttgcag | cttcatcatt | taggccaagc | actgtcaaat | ctatggaatc | ttatggacac | 120 |
| atcatacagt | gagcgacaat | ctttttccca | tggtaatcaa | ttgttgtcac | tctcatctgc | 180 |
| agaagtaaca | gatccgggaa | gtcttactct | agaaatagtc | cagcaggtag | gtatctaact | 240 |
| taaatgatcc | atagaaaata | tcggaaactc | aaatatttaa | aaagtctaca | ac | 292 |
| <210> <211> <212> <213> <400> | 20150 429 DNA Glycine max 20150 | | | | | |
| tatgtgaagt | gaccaaatga | agaatgaaaa | tgagttgttc | gatggtgcat | gtgggggttt | 60 |
| tgtgctgcct | atagatgcag | gaatcgtagt | taaatgggtt | ttaagctact | gaagtgaact | 120 |
| atgtggatat | ggcttcttgt | gcattagaaa | tagccaagta | aaattctgaa | gaactactta | 180 |
| ctaccttggt | attgagattg | aggggttgtt | ctgttcccag | cttccttctt | ccttccgtta | 240 |
| tcaattattt | tattcaactc | tatcgcacct | tgctaaaatt | cttaagctta | ggctgttgaa | 300 |
| tttccttttg | ttcaattata | gaaagagaag | gaaaatatat | tgatttgact | cctgagaaac | 360 |
| aggggttgta | ccggatcaag | aaaaatgttt | tggaatcgga | tggttgtttg | ttgtgcctaa | 420 |
| ttgtctata | | | | • | | 429 |
| <210> <211> <212> <213> | 20151 365 DNA Glycine max | | | | | |
| <223> <400> | unsure at a 20151 | all n locat | ions | | | |
| A. San | tatcatttag | | | | | 60 120 |
| | ctcccgtttt | | | | | |
| | gttctcacat | | | | | 180 |
| aaagagctta | aattggtggg | catcaagaac | caatttcctt | atagaagaga | atgcgcccat | 240 |

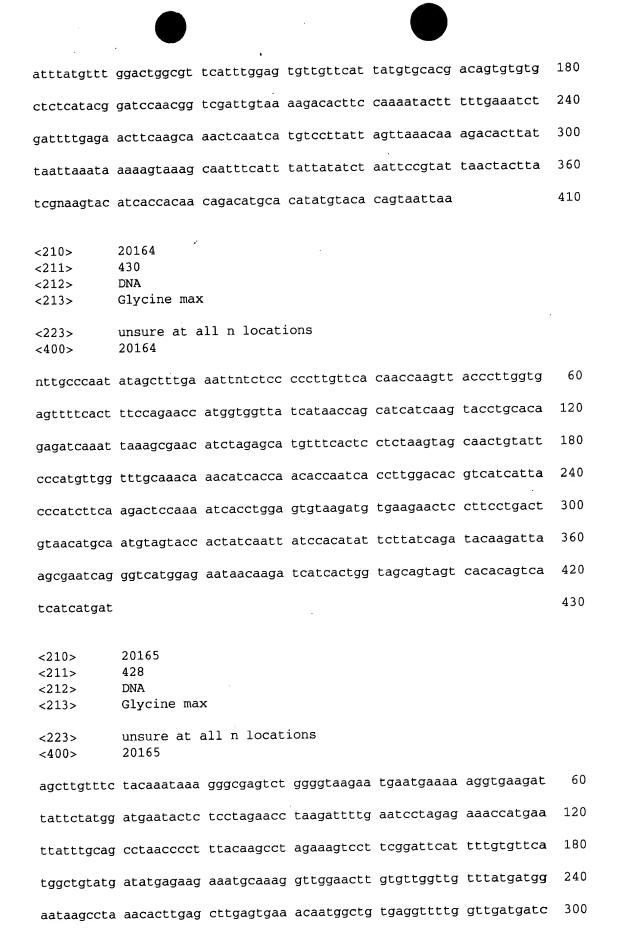
| tattccaaca | cccgtgaaga | ccaccataat | tgaggtgtta | atccaataag | tgaaggatga | 300 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| atttggaggc | ttgtatgtca | tgttgtacat | aagcataggc | agaacgaaat | ccaaagggat | 360 |
| gaaac | | | | | | 365 |
| <212> | 20152 434 DNA Glycine max | ς | | | | |
| <223> <400> | unsure at a 20152 | all n locati | ions | | | |
| ntccaagatg | gccaagttca | agcctaacga | tgggttattt | attgaaaggt | gcaatacaat | 60 |
| tctgtgtatg | acattcaaat | tttactatgt | tagtaggtaa | gattaaggat | tgggcagtgt | 120 |
| tgtcatggac | tatcgaaaag | atttccaagt | aacaagaagc | atccgatgat | gttcgataga | 180 |
| attcatatat | tgttgttgct | ttagaaatat | atttgttatt | taattccaat | tatattttga | 240 |
| ccaatttcat | tttacgtttt | taacaaaact | ctttataatt | ntagtctgct | ataatatacg | 300 |
| aatcattgta | gagtaatatc | ttaaaatttt | aatgattaac | taagatcagt | gtattatgca | 360 |
| tttcaattaa | tataaaaact | tgtatatctc | atagtatata | aaaacttgca | tacttcatag | 420 |
| gaaatatttg | ataa | | | | | 434 |
| <210> <211> <212> <213> | 20153 414 DNA Glycine max | × | | | | |
| <223> <400> | unsure at a | all n locat | ions | | | |
| agcttatcat | gtataaacta | tntaaaatcg | taaggaataa | ctttttataa | gattgaactg | 60 |
| aaaataaata | aattttggtg | gataaaaatc | gtaacaaatt | gtgaaatcgt | aaactcaatg | 120 |
| aacaaagagg | gactaaaagt | aacttttcga | aaatttgagg | gactaataaa | aataattttt | 180 |
| tttgagaact | aaaaaatact | taccgaaatt | tgaaagatta | aaaatatatt | taagccttaa | 240 |
| atcaatctta | taaaatcagc | ttgtaagatg | aaagatgtcc | cacacttata | tacactaatt | 300 |
| ngactatatc | tctagacaat | gtgatctcga | acacaacctc | tcatgtcaaa | gataggatat | 360 |
| ctcatacata | aagtttgcac | gattggaagt | tatgggtagt | gtgatagatg | tccg | 414 |

| <210> <211> <212> <213> | 20154 438 DNA Glycine max | |
|----------------------------------|--|-----|
| <400> | 20154 | |
| actcagcttc | cttaggaacc ttgaaggata tatatatttc tttgcttggt tccttgtatt | 60 |
| acaacatact | ttcatttcca tttgatttgc accactccac ctttcacaca gtaggtaaaa 1 | 120 |
| cctttggagt | ccatggctcc catcgagatt acattcttct taagctatag aacatgaaaa 1 | 180 |
| cattggttaa | agttataaca attccatcat gcattttgat ttgtaagaac ctatgccaac 2 | 240 |
| taccttgcaa | gcggcattgt atccattaga gcaataccac taaattattc tcataggtct 3 | 300 |
| tgaactagct | tttgtgtgga cacatatgat atgaacaacc tgagtcatcc attatccaaa 3 | 360 |
| atactgttgt | tgttgatcaa caatcgagag taccaaatta gttttttgat gaggaattgt 4 | 120 |
| tggatcaagt | ggcctcag 4 | 138 |
| <210> <211>. <212> <213> | 20155 309 DNA Glycine max | |
| <223> <400> | unsure at all n locations 20155 | |
| agcttgtctc | atttagatcc aggaaagaca aggcggttga aggaaccagt tctgctcccg | 60 |
| aatatgacag | ccatcatttt atgagcattg accaccagca acgcttcgaa gccatcaaag 1 | 120 |
| gatggtcatt | ccttcgggag agacatgtcc agctcangga cgacgagtat actgacttcc 1 | 180 |
| aggaagagat | agttcgccgg cggtgggcat cgctggttac ccccatggcc aagttcgacc 2 | 240 |
| cagacgtagt | cctcgagttt tatgccaatg ctnggcctac agaggagggt gtgcgagata | 300 |
| tgcgttctt | · | 309 |
| <210> <211> <212> <213> | 20156 423 DNA Glycine max | |
| <223> <400> | unsure at all n locations 20156 | |





| <210> <211> <212> | 20161 268 DNA | | | | | |
|-------------------------|------------------------------------|-------------|------------|------------|--------------------|-----|
| <213> | Glycine max | (| | | | |
| <400> | 20161 | | | | | |
| tatctcttca | aatcattttg | aaaaggcacg | aactacctat | atatatgggg | gtctgatttc | 60 |
| caaaagcaag | agagagatat | tcccagagaa | cttcattgtc | aaatgctctc | tcaacaactt | 120 |
| ttgggcgaac | acttgccaat | ctattaagag | ttcatccaag | aacttcaaat | gtaatatcct | 180 |
| tcttttaaag | agagaattct | tcttcttctt | attaaaagag | attgattaat | ggaccgagag | 240 |
| tctcttaagt | tgtaaggatt | cctgaaca | | | | 268 |
| <210> <211> <212> <213> | 20162 431 DNA Glycine max | ĸ | | | | |
| <400> | 20162 | | | | | |
| tgataaatct | atatatggtt | taaaacaagc | ctcctgtctg | tggtacctta | agtttcatgg | 60 |
| gataatttct | tcatttggtt | ttgatgaaaa | ccccatggat | caatgcatat | accacaaggt | 120 |
| cagtgggagt | aaaatatgtt | ttctttttt | atatgtagat | gatattttac | ttgtagccaa | 180 |
| tgatcaagtt | ttgctacatg | aggtgaaaca | atttctctct | aagaattttg | aaatgaagga | 240 |
| tatgggtgat | gcatcttatg | tcatcgacat | taagattcat | agagatagat | ctcgaggtat | 300 |
| tttgggtcta | tcacaggaaa | cttatattaa | caaaattcta | gagagatttc | agatgaaaga | 360 |
| ttgttcacca | aatgttgctc | tcattgtgaa | gggtgatagg | tttaatttga | accaatgtc c | 420 |
| aaagaatgac | t | | | | | 431 |
| <210> <211> <212> <213> | 20163 410 DNA Glycine ma | x | | | | |
| <223> <400> | unsure at 20163 | all n locat | ions | | | |
| agctttatag | cataaacgag | attntgcaag | aaattaaaat | tgcaaaaaga | gtttgatcct | 60 |
| attttaatta | atgacatata | atgtttattg | tataatagaa | cctaaaatgt | taattactta | 120 |



| cttccttgat | ttttgtcatg | cttactagct | tatttcagct | gtgattctaa | tgcttatgct | 360 |
|----------------------------------|------------------------------------|-------------|------------|--------------|------------|-----|
| cctatctttg | aaaattgcat | gcttgtgaga | agtaattgat | ntaagcattc | catggtattc | 420 |
| agttcata | | | | | | 428 |
| <210> <211> <212> <213> | 20166 426 DNA Glycine max | × | | | | |
| <400> | 20166 | | | | | |
| tggactcaaa | gagaaactta | gaatggctct | agagtattag | taaaaaaact | ataaaataaa | 60 |
| gactcaacaa | acctctagct | ttggcacttg | ttttcacagt | aattttcaat | tgaaatttcg | 120 |
| gaactaagat | tggtataaca | taggcaccaa | ttatagaata | aattttgagc | caaaacaaca | 180 |
| agcgcacttc | cctttcactt | ttttttcct | ggatactgat | ttttctgcca | acttgtgtga | 240 |
| tttttagtat | tttttcctgt | tatccaaatc | acttggttct | ttttttataa | cttttttcca | 300 |
| gatgtctagc | aaattcagta | aaactttcag | ctcaaaattc | gaagtaacca | attctcagta | 360 |
| atttttacaa | gtttgtatgt | ccaagctgcc | agcaccaacg | attttttt | aagcatggta | 420 |
| tattga | | | | | | 426 |
| <210><211><212><213> | 20167 369 DNA Glycine ma | × | | | | |
| <223> <400> | unsure at 20167 | all n locat | ions | | | |
| agcttcgtgc | ttaaatatgt | atggcaaaac | ttcattactg | ttgttcaaga | catagaagtg | 60 |
| agcttgtaac | aaatcttcta | cacttggagt | gatcacctgo | agtectettg | aacccttacc | 120 |
| acccactctg | tcatcatgco | gacactcang | aagcccaaca | actttagcct | tctctaagta | 180 |
| ttctgaacaa | aattcaatgg | cttcttctgc | aatgtaccto | : tcaacaatag | atgetteegg | 240 |
| acgatataga | ttctttgtat | accetttaa | gatcttcatg | , tatcgctcaa | ccgggtacat | 300 |
| ccaccgtaga | taaacaggad | cacaacattt | gatttctctt | gaccagatgo | acatcaagtg | 360 |
| aatcatgat | | | | | | 369 |

| <211> <212> | 20168 421 DNA Glycine max | | | | | |
|--|---|--|--|---|--|--------------------------|
| | unsure at a 20168 | ll n locati | ons. | | | |
| tccattgttg | agtttttgct | tcccttttca | tgctttattt | cactccccac | aagtaagtgc | 60 |
| aatttccctt | ggttatttgg | ctctccattg | atgtgttttg | gtgctttagt | tgctcatttt | 120 |
| ttgcaaaatt | cgtgaagcaa | tttgcatcta | aatccatgct | tgttttgtgg | agttgaggat | 180 |
| ttgaatgaga | aggccttagg | cctatgttgt | attctgaagc | aatggggcat | gccacattgc | 240 |
| ccccattctc | ttgcaattta | tgtccaaaca | tgtgcccatc | aagtgctcgg | tgaaatgccc | 300 |
| caatgatata | tgaatatgat | tttgcaaaat | tgggatggtg | gggctgtttt | gtgtatgtag | 360 |
| agacagcata | ggaaagtcga | aatagatgcc | caaatgcaat | cccaagctta | ngaacccaaa | 420 |
| С | | | | | | 421 |
| <210> <211> <212> | 20169 360 DNA | | | | | |
| <213> <223> <400> | Glycine max unsure at a | | ions | | | |
| <223> <400> | Glycine max unsure at a 20169 | all n locat: | | tgccctaact | ccttaaggcc | 60 |
| <223> <400> agctttgcga | Glycine max unsure at a 20169 aaagcttgcc | ll n locat | acccatcaac | | | 60 120 |
| <223> <400> agctttgcga ggtgacatct | Glycine max unsure at a 20169 aaagcttgcc atgcccttaa | gctagagctg | acccatcaac atagaatctt | tttccgattt | gatttgtccc | |
| <223> <400> agctttgcga ggtgacatct catgatttac | Glycine max unsure at a 20169 aaagcttgcc | gctagagctg tcttgacttg gtgaatcaag | acccatcaac atagaatctt gctcttatat | tttccgattt | gatttgtccc | 120 |
| <223> <400> agctttgcga ggtgacatct catgatttac atggaacgaa | Glycine max unsure at a 20169 aaagcttgcc atgcccttaa ctaaaatggg | gctagagctg tcttgacttg gtgaatcaag acaagtggta | acccatcaac atagaatctt gctcttatat gtttacatat | tttccgattt gaatgatgca acgagaccga | gatttgtccc atgcacatgc aaaagatcca | 120 180 |
| <223> <400> agctttgcga ggtgacatct catgatttac atggaacgaa tctttntaat | Glycine max unsure at a 20169 aaagcttgcc atgcccttaa ctaaaatggg aagaagaaaa actacgctct | gctagagctg tcttgacttg gtgaatcaag acaagtggta angcattgcg | acccatcaac atagaatctt gctcttatat gtttacatat gcgccctaac | tttccgattt gaatgatgca acgagaccga gtatgcatta | gatttgtccc atgcacatgc aaaagatcca canagtgacg | 120 180 240 |
| <223> <400> agctttgcga ggtgacatct catgatttac atggaacgaa tctttntaat | Glycine max unsure at a 20169 aaagcttgcc atgcccttaa ctaaaatggg aagaagaaaa | gctagagctg tcttgacttg gtgaatcaag acaagtggta angcattgcg atatcactag | acccatcaac atagaatctt gctcttatat gtttacatat gcgccctaac | tttccgattt gaatgatgca acgagaccga gtatgcatta | gatttgtccc atgcacatgc aaaagatcca canagtgacg | 120 180 240 300 |
| <223> <400> agctttgcga ggtgacatct catgatttac atggaacgaa tctttntaat cgttactcta <210> <211> <212> | Glycine max unsure at a 20169 aaagcttgcc atgcccttaa ctaaaatggg aagaagaaaa actacgctct aagagacaag 20170 425 DNA | gctagagctg tcttgacttg gtgaatcaag acaagtggta angcattgcg atatcactag | acccatcaac atagaatctt gctcttatat gtttacatat gcgccctaac | tttccgattt gaatgatgca acgagaccga gtatgcatta | gatttgtccc atgcacatgc aaaagatcca canagtgacg | 120 180 240 300 |

| ctgagcaact | ccttcttcaa | tatttctatg | agggacttag | caacatggag | aagagtatga | 120 |
|-------------------------|-----------------------------------|-------------|--------------|------------|------------|-----|
| ttgatgctgc | tagtggtgga | gctcttggtg | atatgacccc | tgctgaggct | aggaatttga | 180 |
| ttaagaagat | ggattccaac | tcccaacaat | tcagtgcaag | aaatgatgct | attgtcctta | 240 |
| gaggagtcca | tgaggtggcc | acggattcat | cttcatctac | tgaaaataaa | aagcttgagg | 300 |
| gaaaacttga | tgccttggtt | aatctagtaa | cttagcttgc | catgaatcag | aaatctgcac | 360 |
| ctgttgcaag | agtctgtggt | ctatgttctt | ctgcagatca | ccatacagat | ctttgtcctt | 420 |
| ctttg | | | | | | 425 |
| <210> <211> <212> <213> | 20171 424 DNA Glycine ma | × | | | | |
| <223> <400> | unsure at a | all n locat | ions | | | • |
| agcttatgat | aatgaagttg | ttacaaattt | tatcttgaat | gaaactgtgc | caaaaataga | 60 |
| tgtcatgaag | ttgttacaat | atataatgca | cttaataacc | ttgcaaataa | acttcagata | 120 |
| atcttaatct | ttggaaatgt | attgatatta | ggacaaatgg | tcatcacata | tgcaacaaac | 180 |
| atatagccat | cctaatttac | tcagttacat | gaagctngtt | atgttcaaaa | tatcacacat | 240 |
| gcaaattgat | gacaccttat | agataatagc | cttagaagtt | gattatcatg | actcanaatt | 300 |
| aagggtttca | ccattacact | atcatatcat | ttaaccacaa | cagagaattt | aatacaaagt | 360 |
| cacactaaac | ttgagttaca | tcacatctac | ttatggcact | aagtataaac | attgcatgaa | 420 |
| gtat | | | | | | 424 |
| <210> <211> <212> <213> | 20172 467 DNA Glycine ma | x | | | | |
| <223> <400> | unsure at 20172 | all n locat | ions | | | |
| gcgatggccn | ttgaagacct | cagtcatacc | ngcgacacta | taaaatacto | acgcttgagg | 60 |
| aacaatgggc | ctaccctaca | cctgagatgn | gtgattcaaa | tgtgtccaat | gccccacgga | 120 |
| +~+++>++ | . daaattaaaa | teettenaaa | . aaaacaatat | ataaattcoc | ccaataaggg | 180 |

| ccaaaatgaa | gtgccacagg | tgctctaaaa | ggtgaaggag | catttgtcag | ccccaaaggg | 240 |
|-------------------------------------|---|-------------------------|------------|------------|--------------------|-----|
| gtgaccaaaa | actcctatat | ggccctgggg | tgccgaaaag | tggtttggtt | aaattaagc <u>c</u> | 300 |
| ccttcatctt | aaatatttgg | tgacctccta | taggcgcact | ttttaaaaca | ccaggccctt | 360 |
| cctattattc | atctactcct | caatggtggg | gtgggggtta | atccttaccg | actcgcggtc | 420 |
| ttatttgacc | attcccccaa | cagcctgtct | cctcctctt | taccacg | | 467 |
| <210> <211> <212> <213> | 20173 318 DNA Glycine max | × | | | | |
| <400> | 20173 | | | | • | |
| agcttgtgtt | ttcaaccctt | atcactgctt | tatgtaaagc | tagagaagtc | acttctgatt | 60 |
| ctaggacact | ggagagtctt | caccccgcca | ttaatttggc | atatgtgaag | aagaactatt | 120 |
| ggaatcttga | tgatctaata | gtgactttca | gagggcctat | gaaggccaac | gggaagaaat | 180 |
| cgaagactct | cccatctttt | gaggttccct | ctaccacatc | agcaccaact | tcttcttacc | 240 |
| taggtacttc | tgctcatcac | caacttcttt | agattttctt | ttacactaag | atgctcatgc | 300 |
| catgatgcaa | gcctacct | | | | | 318 |
| <210> <211> <212> <213> <223> <400> | 20174 434 DNA Glycine ma unsure at 20174 | x all n locat | ions | | | |
| ccgctntttg | gagtagaaac | atgggaccaa | cacattttat | tttttaaagt | cgtatctagt | 60 |
| caagatctga | gagaccatac | aagtttccta | gcggtttcta | attatatggg | ccattaagtc | 120 |
| tatcatatgc | tgacaatagc | cgagaagccc | atgaatttct | teggggeegg | agtaggtgtc | 180 |
| tgccattgcc | ttggccttgg | ctaataatcg | aggaagttct | tgactcccgt | tcaaggtaag | 240 |
| agcaaaccgg | tccatccaca | tggttgcctc | ttggtgtaaa | gagtcgatca | cccttcctct | 300 |
| agcctctttt | ttcgcgtata | ctagggcata | ctcgtccgcg | accctatgct | cgtgggccgt | 360 |
| gactagactt | aactcttctt | ggtacttggc | aatgatagct | agcatgttgg | tctccgtctc | 420 |

| gcataaacgc | tgag | | | | | 434 |
|-------------------------|------------------------------------|-------------------|------------|------------|------------|-----|
| <210> <211> <212> <213> | 20175 193 DNA Glycine max | . * | | | | |
| <400> | 20175 | | | | | |
| catggcggtg | caaagctgac | agtctctgtc | tggagagaga | aagagatacg | cggggaatgt | 60 |
| tgctcgcttt | gaggatggag | ttccgcgcga | tctcggttct | tcatgtgcgc | acacgtggca | 120 |
| agttgccctt | ccccacgctg | agattgaatt | ctcactgtgt | acatgtatca | tagggtccct | 180 |
| cacaccttgc | caa | | | | | 193 |
| <210> <211> <212> <213> | 20176 566 DNA Glycine max | ¢ | | | · | |
| <223> <400> | unsure at a 20176 | all n locati | ions | | | |
| tgtacacgcg | atacacgtat | ggtatggtaa | ttacgattat | cgcgatatat | aagagaantg | 60 |
| tgacgacaca | canacccaca | agaggagggc | attgatgacc | tcgatagaca | cgcccgctan | 120 |
| nacaaanaan | accccagcat | gcatgaaaag | tggcaaagca | aacaaacact | ttcttctaac | 180 |
| aggcatcacc | agccgaagtg | agagaaaaca | aacacttgca | atgaacattc | acggcaatgg | 240 |
| gcaaaagcca | gaacataatg | ataacgactt | acagggagca | aaataagtga | actgaccgct | 300 |
| gctgatgatt | agaaatcggc | atataagatg | gataccagcg | cgcaacaatt | cactgcacgc | 360 |
| aatgaagcaa | gaatacctag | atgagacctc | gaggagtgga | cggaatagtc | gaaacgccac | 420 |
| acaactaagg | agctgagaga | aaacgtgatg | gacatgctaa | gaaagacacc | gaacatgaaa | 480 |
| tgaatcacat | accatatacc | agaaccacag | acgaagcacc | ataagactaa | acacatcacc | 540 |
| aaaagagcga | cgccgtgcta | caagac | | | | 566 |
| <210> <211> <212> <213> | 20177 170 DNA Glycine max | x all n locat: | ions | | | |

| <400> | 20177 | | | | | |
|----------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| ggtggttatg | gctcgccaat | ggaggtggtg | gtgctgaagc | gttgangagg | taaaggctga | 60 |
| aaggttgcgg | aacgagatat | ctcatggata | cacatattgc | ttaggcaatg | cctggacgtc | 120 |
| cactgatgag | acaggaatag | taacggtgac | agataggaac | tgatctacgt | | 170 |
| <210> <211> <212> <213> | 20178 504 DNA Glycine max | × | | | | |
| <223> <400> | unsure at a | all n locat: | ions | | | |
| ggaatcacag | cagcaacaan | tatgtaatag | aacaaacaac | cnccccaggg | attgaacttg | 60 |
| aacaccacca | aanaccaact | ccagaactca | aagcttttat | agacacacca | aatctttgga | 120 |
| gcaaagcagg | atacaatcag | gacattcctc | ctcctgaaac | tcctatcgac | cactcgctat | 180 |
| cataagagga | gaatcaccgc | cctttcaatt | tatgcaatct | atactggggc | atacccgata | 240 |
| ccctatttta | taagacagga | ccaaacttcc | gaagttatgc | aagaaattcc | tctcttagca | 300 |
| ttactaggaa | cacagtaata | gaaattggcc | atcacaaagc | cttcgtagca | accttagcat | 360 |
| ggacaatacg | tctatatgta | caataaaagc | gcacactcgg | aatcatagaa | acaaaattac | 420 |
| tcaactgtac | aacaatacga | ccgcagaatc | agaagcgtca | tacctaacac | acgaagacct | 480 |
| ctcctcaact | gcatacacaa | accn | | | | 504 |
| <210> <211> <212> <213> | 20179 391 DNA Glycine ma | x | | | | |
| <223> <400> | unsure at 20179 | all n locat | ions | | | |
| tggtgagctc | : tgatatcgca | cttggaaggg | gaatatctat | caatacgacc | ttataaaata | 60 |
| tcacatgaag | cccaaaacac | attatgtcat | attatcagga | ggttttaagg | aatacctgga | 120 |
| tccgtagaga | gctngatcaa | . ctcgcagcgg | aatctaagag | taggcacaga | caatttggat | 180 |
| taatgacctt | gctcaaccaa | gtcacattct | tctcttatga | gaccttcgct | ttctcttcag | 240 |
| atggtgagag | gacgaattgo | : ttattgagtt | tgtgtattgg | ggaccataac | catatcttat | 300 |

| gtggtaaacc | tattatggta | ccattatccc | ttagtggacc | acactgattc | tcgctcatta | 360 |
|--|---|---|---|---|---|---------------------------------|
| agttcatatc | cattcttctg | gcagtctaat | g | | | 391 |
| <210> <211> <212> <213> | 20180 434 DNA Glycine max | ς | | | | |
| <400> | 20180 | | | | | |
| cgtaggatta | tggggtaccc | atcacatgtg | gtactaggtg | tcggtcgggc | gatggtgcac | 60 |
| atcaagtttt | ccacatccac | aaagcgcgca | taaacccacc | atcccctgtt | gcccacctcc | 120 |
| aactgagctc | acgtagccca | tatcctcgtt | tctctcaaca | ccgggtcccc | atcaatcctc | 180 |
| tcaagcttcc | acaacatcca | agcaaaacaa | cattcaaaca | gcacaagcta | tcacagccaa | 240 |
| gcaaaacaga | gcaaaggcag | aaaactctgc | cacaacacca | accaaatcac | agcttttctc | 300 |
| acttaaagac | cccagtaaca | attcctacga | tccaattcgt | taaccgctgg | atcgactcca | 360 |
| aaattttact | ggaagtatat | agtacatgag | cctacattgt | gaccgttggg | atctactagc | 420 |
| aaacatccag | aact | | | | | 434 |
| | | | | | | |
| <210> <211> <212> <213> | 20181 568 DNA Glycine ma: | x | | | | |
| <211> <212> | 568 DNA Glycine ma | x all n locat: | ions | | | |
| <211> <212> <213> <223> <400> | 568 DNA Glycine max | all n locat: | | cacgatacaa | ccccacnncc | 60 |
| <211> <212> <213> <213> <223> <400> | 568 DNA Glycine mas unsure at a 20181 | all n locat: atcggacaca | gccacggccc | | | 60 |
| <211> <212> <213> <223> <400> ccccctcac | DNA Glycine max unsure at a 20181 ccngcagagg | all n locat: atcggacaca tgatgccatg | gccacggccc | cgaataaact | cggcccgcgg | |
| <211> <212> <213> <213> <223> <400> cccccctcac caacaacaga gaacctcaag | DNA Glycine max unsure at a 20181 ccngcagagg gcggtnngnt | all n locat: atcggacaca tgatgccatg caggcgcgca | gccacggccc aacctcacgg atcttttta | cgaataaact | cggcccgcgg | 120 |
| <211> <212> <213> <223> <400> ccccctcac caacaacaga gaacctcaag agacatcgtc | DNA Glycine man unsure at a 20181 ccngcagagg gcggtnngnt agtccacccg | all n locati atcggacaca tgatgccatg caggcgcgca caacgggcac | gccacggccc aacctcacgg atcttttta gtaacggaaa | cgaataaact caacgctcca aagagcctct | cggcccgcgg tacacactcg aaacttgcac | 120 180 |
| <211> <212> <213> <223> <400> ccccctcac caacaacaga gaacctcaag agacatcgtc acccaagtct | DNA Glycine ma: unsure at a 20181 ccngcagagg gcggtnngnt agtccacccg tcaaagatcc | all n locat: atcggacaca tgatgccatg caggcgcgca caacgggcac aacagtgaac | gccacggccc aacctcacgg atcttttta gtaacggaaa aaacgctggg | cgaataaact caacgctcca aagagcctct catccacact | cggcccgcgg tacacactcg aaacttgcac attgaggcag | 120 180 240 |
| <211> <212> <213> <213> <223> <400> cccccctcac caacaacaga gaacctcaag agacatcgtc acccaagtct ccccacgtag | DNA Glycine max unsure at a 20181 ccngcagagg gcggtnngnt agtccacccg tcaaagatcc gagaagagcc | all n locat: atcggacaca tgatgccatg caggcgcgca caacgggcac aacagtgaac agagcctata | gccacggccc aacctcacgg atcttttta gtaacggaaa aaacgctggg taaaggctct | cgaataaact caacgctcca aagagcctct catccacact ctccacaagc | cggcccgcgg tacacactcg aaacttgcac attgaggcag ttcctcgtgg | 120 180 240 300 |
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| cagccatctc | acggtgcgcc | ggaccgcc | | | • | 300 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| <210> <211> <212> <213> | 20182 386 DNA Glycine max | ς | | | | |
| <400> | 20182 | | | | | |
| tatagcaggt | ccacatgctc | gccgattgtt | tgtctgacat | atacaatggc | tcaaactaat | 60 |
| acgaggatac | acatactcca | tcacaagtgc | tgcatcgtca | atgtcaaata | tgatgctcac | 120 |
| aacatgtcgt | tgttaacata | gagataccag | agtggcatta | ccatgaactc | ataaatgaca | 180 |
| tggactgttc | cagatgctgc | atgatgaaca | tctacactac | tcatgcacat | aactagtaaa | 240 |
| acgtacataa | tgtactcata | ccataactat | actagcaaat | cctaatgcac | ataacatcat | 300 |
| atcgattgta | agaatcgtat | aattgttcct | gactgtcctg | tgtctgaacg | cactactatc | 360 |
| tacacaaaag | agctacagag | accatc | | | | 386 |
| <210> <211> <212> <213> | 20183 575 DNA Glycine max | ĸ | | | | |
| <223> <400> | unsure at a 20183 | all n locat: | ions | | | |
| gagatgcnnn | tttaatnccc | gtaagganca | ncngacacnt | atataaaatc | acncgcgtcg | 60 |
| tacacctatg | agctatattg | tgtgtggaga | agatttatcg | ataatcaagc | gttatatccc | 120 |
| acctaggaac | aatgaggcgt | gtactaccat | cccgctattt | catcccttcc | gcactatccg | 180 |
| acctcgcata | gtcacatgcg | taggaaccat | acacctatta | aatgcgtacg | atgtatgagg | 240 |
| aactctccac | atggatacca | cgttctcatc | atatacaaga | ctatcatcat | tgctnagaca | 300 |
| ttaactatca | ggctcatgac | ctgtccagtt | actcgataag | gaccactact | ctttgagaag | 360 |
| acgaaatgcc | taataccata | atagatctcg | accctcaaac | cgagtgaggc | gcggctccct | 420 |
| acaccgaatt | gttcaaacct | aagagatggc | tcgaacatct | cactatcatt | atctgactca | 480 |
| gaaacagaca | cacgcaagac | taacaagtat | gctctattga | tagctccaaa | caaacatgtc | 540 |
| tgaatcagct | aatattgact | gcgtctagta | taacq | | | 575 |

| <210> <211> <212> <213> | 20184 377 DNA Glycine max | ς | | | | |
|----------------------------------|------------------------------------|-------------|------------|------------|------------|-----|
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| atgcttcttt | cactaattta | taactggaga | atatatttat | gtttaacaaa | actagttcat | 60 |
| aagttatata | aaagtttact | aatttttgaa | aatţtattac | aatgacttat | tcaattataa | 120 |
| tagttgatta | aaattgggaa | ttgaagtagt | gtaagataat | aaaaatggac | atatatttt | 180 |
| acataaattc | tagagtaaaa | tatgttttta | gtccttaaaa | aaatttacaa | atttgatttt | 240 |
| agtcattaaa | caatcttatt | tttgtccccc | taatatagaa | ataataatgt | cacaatatat | 300 |
| actatcaaga | tcgaagatag | agtatttcca | atttagagga | gcaaatacag | acaaaagaat | 360 |
| ttagatgact | ctaaatc | | | | | 377 |
| <210> <211> <212> <213> | 20185 233 DNA Glycine max | x | | | | |
| <400> | 20185 | | | | | |
| ccactatctg | cacgccaaga | ctatatctcc | atccacagac | ggaactcaaa | aaccccaacg | 60 |
| gcggagaagc | gcggaaatga | attccagacc | atgagtccaa | atgtaaagac | aagccaacgg | 120 |
| caaaaaaggc | caacatcgta | acaacatagg | gacagatctg | agcgtatgct | ggaaaagaga | 180 |
| aagaacagag | cgagggataa | gacatccact | acaaacataa | actgaagaag | ccc | 233 |
| <210> <211> <212> <213> | 20186 574 DNA Glycine ma | x | | | | |
| <223> <400> | unsure at 20186 | all n locat | ions | | | |
| cgaccccgcc | acacgcaaca | gcgtaacggc | ggactacgcg | aacaacaaga | gaacttcccn | 60 |
| ccncancnac | aggagggtna | nnttgaagcc | gtngaacacc | caaggcgaan | acgagcacgg | 120 |
| cacccggaga | tccacnagag | acgaccggca | agcacgcaag | cttagcatga | aaggaggcgc | 180 |
| gaagcgccaa | caacaagcta | acctgaagaa | caccgcacag | gagcacagca | cacgaaaagg | 240 |

| cgcgaagcct | gaaacactca | ccgcaacgcg | aagctagatg | cacgcgaaac | ctcgcatctc | 300 |
|-------------------------|------------------------------------|-------------------|------------|------------|------------|-----|
| aagcaaagcg | cagaatgcag | aaagacaata | ggtgaagcag | aaccactacc | caagaggcta | 360 |
| gggaaagata | cacagatgtc | atagagccca | agaagccttc | ggcacagagg | aacgcgacgg | 420 |
| acgaccacac | acaacctgaa | gagaacaatg | aggaaccgcg | agacacacca | ctangacgaa | 480 |
| gaggcgacag | acaatggaga | cagccacatc | gacgatagca | acgcgaagga | cgccacgccg | 540 |
| accgcgcaga | acccgaagac | agaacaccaa | accg | | | 574 |
| | 20187 523 DNA Glycine max | k all n locat: | ions | | | |
| <400> | 20187 | | | | | |
| acgccccagc | gagaaaagaa | cgacggaagg | aaaaacacac | acccagagac | ttgagccctg | 60 |
| accccccnaa | naaaaacaac | accgaaaaac | caagaaacca | ccgcaaccnt | aatttcagac | 120 |
| cagaaaggga | aaccaggggc | ggagcaaaaa | ggacccacaa | acagagcagg | ggaacacaga | 180 |
| cacggcccac | ggaacacaca | cagccgcggg | agacgaaccc | acagaagaaa | cgcacggcac | 240 |
| aaacgccaca | gccgccgcaa | acagaaaggg | ccgcatccaa | atacaaacac | ccgaacaaaa | 300 |
| aacgcgaaag | aaccgccgaa | gagcaccaca | gatcgagcca | acgacaaaag | cegeggcaaa | 360 |
| aacaaagccc | aaaaaagcac | ggcgaacact | cggcggacag | acagaaggag | cacgcgcaga | 420 |
| caccgacagc | gaccaccaat | aagcagacca | aggacacacg | accctaagaa | gggacaaacg | 480 |
| gcacaagaac | ccgacgcacg | aacacaacga | gacaagcagg | aat | | 523 |
| <210> <211> <212> <213> | 20188 449 DNA Glycine ma | x | | | | |
| <223> <400> | unsure at 20188 | all n locat | ions | | | |
| attcagctcg | gacccgggat | cctctgagtc | acctgcggct | gcaattttt | tttgggtaga | 60 |
| tgaagatgta | aaggaggatg | aggatgtaga | agaggaagag | aaatgtaggg | atgagactca | 120 |
| gaagcaactt | gaagaggcaa | gaaaagaaaa | tcataaacta | cataagaagc | ttgatgttaa | 180 |

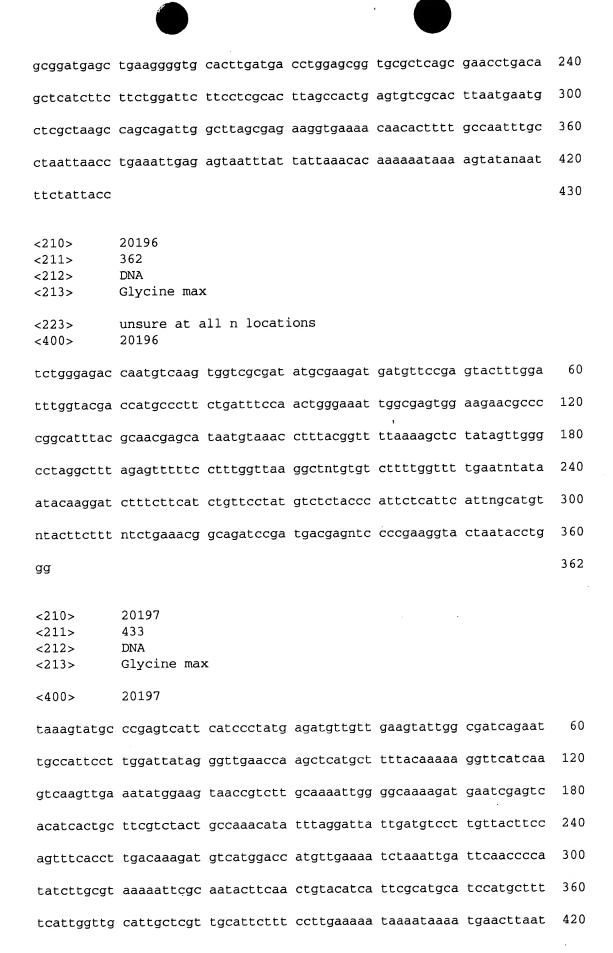
| gagaagaaaa | agatcgattc | tgctttgtgt | cgtacgactg | tgtttggtaa | tcacattatt | 240 |
|----------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| ctcctgtgtt | gtgcgcgtgt | ggaactaagt | agggagttcc | ataatgtgag | gatataccaa | 300 |
| gggaaggaag | tctattggat | tacgatttaa | aaaaaaatta | taaaaaaaat | tgtgttattt | 360 |
| aatcagaatt | tttaaatgtt | ataaataacg | tctgtgatat | tcaattacaa | tntttaagat | 420 |
| tttctacaga | agacataaat | ccgatgata | | | | 449 |
| <210> <211> <212> <213> | 20189 432 DNA Glycine max | ς | | | | |
| <223> <400> | unsure at a 20189 | all n locati | lons | | | |
| tcgtgacgaa | ctatttatgg | aaaaacttca | ttgtngttat | ttagtatata | caaatgagtt | 60 |
| tgttgcaatt | tctctaacat | gcaaccctcg | tgaacccttt | cctcccactc | tctcatcatg | 120 |
| ctgagacttg | ggaagcccaa | aaggttccac | cttttcaatg | tactttaaac | aaaatttaat | 180 |
| agcttctttt | gcaatgtacc | tttcaacaat | ggatgcttca | agatggtata | tattcttcgt | 240 |
| ataccctttt | aagatcttca | tttatcgctc | aaccggacac | atccatcgta | aataaatagg | 300 |
| atcacacaat | tgaacttccc | ttaccagatg | aacaattaag | tgaaccatga | tgtccaaaaa | 360 |
| tgaaggagga | aaatacatct | ccagctaaca | taagataata | gcagtctcat | tttccaagtc | 420 |
| atctaacttt | at | | | | | 432 |
| <210> <211> <212> <213> | 20190 402 DNA Glycine max | x | | | | |
| <400> | 20190 | | | | | |
| ttgcttgtta | aatatgtcta | tcgcatttcc | ctcgtggtgt | aaaatgttac | attgttctgc | 60 |
| tgaataattg | caaaaagggg | ctatgttatt | tgtccagttt | ttattttctg | aatgtctttt | 120 |
| ggggttgata | ataatgctta | gaatttacgt | tttggatcta | attcaatccg | aaagttaact | 180 |
| tgtgaggacc | tattttttgg | ctatatcgct | agttaataca | gaaatctcaa | cacactcttc | 240 |
| ttatgttaga | aataaatttg | atcaagtact | atgactatga | gtattcaaga | ccgaaacata | 300 |

| gaaatgacca | atgcaattta | gtaggaaaac | ctcttatgcg | tctgaagtta | atttctaatg | 360 |
|-------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| taatacaatg | tttcaattca | acattggaat | accataattg | at | | 402 |
| <210> <211> <212> <213> | 20191 425 DNA Glycine max | × | | | | |
| <223> <400> | unsure at a 20191 | all n locat: | ions | | | |
| ttcaacgccg | gccacttcat | atatcnttcc | cattgctacc | aagagtgtac | aatgatcacg | 60 |
| ttgcacatac | aaccccatag | tcggtatcac | attagcaaca | cgttcttccc | ttgacaatga | 120 |
| ggtcacaaca | gtttcagcat | tggtctcctc | tgggctcaca | cgtgcaccta | attggtgtat | 180 |
| atctatatct | gcctcagttg | ggggagagta | ctgcgatccc | ctttgggata | actccgtttt | 240 |
| gatggcctcc | ttcaactcct | atttcattct | ctctagactc | cattttgttt | tcttcttcca | 300 |
| gctcattcct | ccattcttcc | tcaagggttc | caattatctc | agccacctgt | tgttgggtga | 360 |
| tggatgtgga | cgagttgctg | gaaccacatg | atgcccttcc | atattattga | ctaattgtga | 420 |
| cacta | | | | | | 425 |
| <210> <211> <212> <213> | 20192 374 DNA Glycine max | . | | | | |
| <223> <400> | unsure at a | all n locati | ions | | | |
| tctagcttgc | ctcanagaga | tccagganag | acaaggcggt | tgaaggaacc | agttctgctc | 60 |
| ccgaatatga | cagccatcat | tttaggagca | ttgaccacca | gcaacgcttc | gaagccatca | 120 |
| aaggatggtc | attccttcgg | gagagacatg | tccagctcag | ggacgacgag | tatactgact | 180 |
| tccaggaaga | gatagttcgc | cggcggtggg | catcgctggt | tacccccatg | gccaagttcg | 240 |
| acccagacgt | agtcctcgag | ttttatgcca | atgcttggcc | tacagaggag | ggtgtgcgag | 300 |
| atatgcgttc | ttgggtgagg | ggtcagtgga | tccctttcga | tgcggatgcc | ctcagccagt | 360 |
| tcttgggata | ccct | | | | | 374 |
| <210> | 20193 | | | | · | |

<211>

419

| <211> | 419 | | | | |
|-------------------------|--|-----|--|--|--|
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| <213> | Glycine max | | | | |
| ·<400> | 20193 | | | | |
| ttgtgtaatc | gattaccctt atttggtaat cgtttatcag tgactgtcta tgataaatca | 60 | | | |
| aaagatgtaa | ctcttcaaaa aggtttttga ctttttcaaa ttggttttaa gtttttctaa | 120 | | | |
| aagttataac | tcttctaaat ggtcttcttg accagacatg aagagtctat aaaagaaagg | 180 | | | |
| ctttgttttg | catttcaatg atcaaaaaca cttattcata caatccttta caagccttga | 240 | | | |
| atctctttga | acttcttatt cttccttgta ccaaaagcta tctgaagttg tctgggtttc | 300 | | | |
| taaaccttga | aaacttgtgc tattcatctt ttcattctct tctccctttg ccaaaaagaa | 360 | | | |
| ttcgccaagg | actaaccgcc tgaatctctt ttgtgtctct cttctccctt ttccaaaag | 419 | | | |
| <210> <211> <212> <213> | 20194 282 DNA Glycine max | | | | |
| <223> <400> | unsure at all n locations 20194 | | | | |
| cgagaatggt | tatatatcta gcacaagaat tgagccacat tctaattgat tggttctata | 60 | | | |
| tattcatcac | aacaatagaa cctgactcaa atatttnttt aacaaatctt cttaatatat | 120 | | | |
| ctctagaagc | tagagcagtg ataataaggg agatgcatca atacatgaca tcaagataga | 180 | | | |
| tagcaaccgt | gatgttgaat aattctacaa agttcactaa tgtgatagga ctntgattgt | 240 | | | |
| gatccctact | aanaaccaag cttaacanaa ggaaactttc tc | 282 | | | |
| <210> <211> <212> <213> | 20195 430 . DNA Glycine max | | | | |
| <223> <400> | unsure at all n locations 20195 | | | | |
| tcaagccaag | gccagactct tgtgcatcca gaggcttctt tgaaaaaatg ccaaactccc | 60 | | | |
| ttcaaaaaat | ctaatttcaa gcttaaatag gtgggttggt ccgtgctcgc gtgcttagca | 120 | | | |
| caaatcttaa | tcgcttagcg cgcataagtg gattttggct tagcgcactt ctctcgctta | 180 | | | |



| cattgttata | aaa | | | | | 433 |
|-------------------------|------------------------------------|----------|------------|------------|------------|-----|
| <210> <211> <212> <213> | 20198 323 DNA Glycine max | | | | | |
| <223> <400> | unsure at all 20198 | n locati | ons | | | |
| atcatcaaaa | cattcagctt gat | cctttgt | ctacattatg | ttgacaccag | agccatcgcc | 60 |
| aactaattac | taatcagtac cat | gataagg | attgttacag | tgtagatttt | tgcacaatag | 120 |
| tccatgatcc | agttattttg caa | acggaata | aggtgtggtc | ttggcatgat | gagtactang | 180 |
| tccacttgga | cattntggat ago | ccatgaag | tttatagcaa | accatggcag | tgttccctag | 240 |
| tttctcatag | tattgacaaa cta | acatttt | tgttatgcct | cgaaatatgg | atctgccata | 300 |
| tgttgaagtg | gtttgttaac att | . | | | | 323 |
| | | | | | | |
| <210> <211> | 20199 427 | | | | | |
| <212> | DNA | | | | | |
| <213> | Glycine max | | | | | |
| <223> <400> | unsure at all 20199 | n locati | ons. | | | |
| tatgttaatg | ggtcttaaga gga | aaagctca | ctaacattct | taaacttact | tattaaacat | 60 |
| gctcatgaaa | actattttt cto | caattaaa | ataaatccct | tttattttcc | tacaacaata | 120 |
| acccaaacta | atatttattt att | ttatttat | ttatttgcaa | tatataatat | attgtgagaa | 180 |
| aattacatta | gcactccctc aaa | atattgac | ttattacaca | gacaccccat | tcataaaatt | 240 |
| gcatttcttt | tgcaccccct tcg | gtgtcttt | gttaacttat | gctacttaaa | agtgaaaggt | 300 |
| caaaattaac | cttctcactt aac | ccagcact | caacacacct | tntgtttatt | gtttatgaaa | 360 |
| tttgaaaaca | acaaatgtgc tct | tactaggg | tttgatagcg | agggggagca | tgttgctcat | 420 |
| gttgttg | | | | | | 427 |
| | · | | | | | |
| <210> | 20200 | | | • | | |
| <211> | 397 | | | | | |
| <212> <213> | DNA Glycine max | | | | | |

| <223> <400> | unsure at all n locations 20200 | | | | | | |
|-------------------------|------------------------------------|------------|------------|------------|--|-----|--|
| agctttatac | ttgagattga | atagtttaca | atctccgtcc | agatctagaa | gtcatagaca | 60 | |
| accaacttaa | gaagaaaaag | aggaaaagga | gtattcatat | ggaacaagtc | atggaagaag | 120 | |
| gcgaagaggt | gaaccaagaa | tagataatca | tttggggagc | attaagatga | caatccctac | 180 | |
| atttcaaggc | aaaaacaatc | ctgagttgta | tttagagtgg | gagtgaaagg | ttgaacatgt | 240 | |
| gtttgattgc | cataattatt | ctgaggaaaa | aaagatttaa | ctagttgttg | ttgaattcct | 300 | |
| tgattatgct | agtatttggt | gggatcaact | tgtgactaat | angcacagaa | atggtgaaag | 360 | |
| gcctattagt | agatgggagg | agatgaagac | tgtcatg | | | 397 | |
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| agctcacctc | cttgagaagc | ttccttaaga | agattcgtaa | agaagctaga | gcttagctac | 120 | |
| acatacctct | ctaatagcta | agctcacctc | cttgagatga | gaagctagag | cttagctaca | 180 | |
| caccccctat | aatagctaag | ctcaccccca | tgacaaaaaa | catgaaaata | aaaaaaaaa | 240 | |
| gtccttatta | caaagacaac | tcaaaatgcc | ccgaaataca | aggctaaaac | cctatactac | 300 | |
| tagaatggcc | aaaatacaag | gccttgacga | aggaaaaacc | tattctaata | tttacaaaga | 360 | |
| taagcgggct | catacttagc | ccatgggctc | gaaatctacc | ctaaggctca | tgagaaccct | 420 | |
| a | | | | | | 421 | |
| <210> <211> <212> <213> | 20202 73 DNA Glycine ma | x | | | | | |
| <400> | 20202 | | | 20005 | #2#2################################## | | |
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| taagtgtaag | cca | | | | | 73 | |

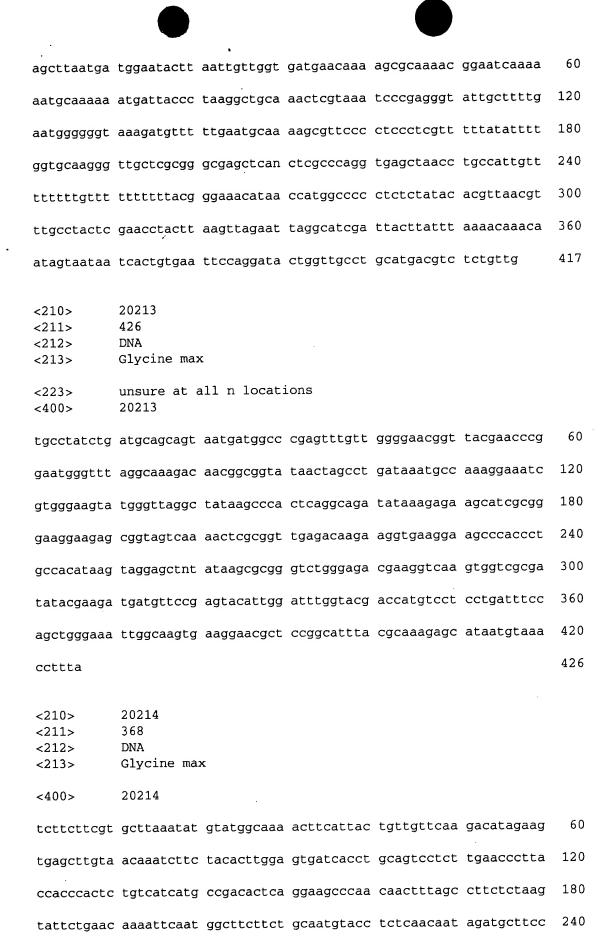
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| gttaccgatt | cttcttcatt | gctatcattg | attgttactt | ttgtcttcat | gacatctgtt | 120 |
| cccactccaa | atgaacaaga | tttagtcgaa | atatttatca | tcccatgtta | caagccttat | 180 |
| catatgctct | atcttaagtt a | aaatgttttt | ctaatgtgtt | ttaactaaaa | aaacaaatga | 240 |
| gaatgaaaat | cgtctaccga | ttgtcttttt | ataaaaacaa | aaagaatgga | atggtttggt | 300 |
| ctaataaata | ttggagagac | ataaaacatg | aaaaaaaacg | taatttttaa | gagaaacagt | 360 |
| gctctgcttc | tcaacatgga | tgaacaacat | attatataat | tagacaaatg | С | 411 |
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| | | | | | | |
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| <400> attttgtatc aaacacaaca tcatgtacca | 20204 tacatagagt atcatagaga | agaagataga cttctcttga gattgtttca | tcattgatcc atccttggct aaccatacaa | ggtgataaag ggacctttgc | ctatcaaacc agttgacaaa | 120 |
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| tgtacataac | tcgtgaaatg | aaatccatgt | ttaggttgat | ttgtttcctg | aatattțacc | 120 |
|----------------------------------|------------------------------------|--------------|------------|------------|------------|-----|
| taaatagatt | gattctaatt | ccataaactt | atttactgta | caattatcat | tgaaatacag | 180 |
| cacacgttat | tcagtccaat | caatggagtg | atttttggtt | tttctagttc | atagtcatag | 240 |
| cagacccatg | cattagcgcc | ccctaaccca | catatactaa | agggataaat | tgatttgggc | 300 |
| cattttgttc | ttttattaag | tgaatggtta | gtgctaggtg | ggttaaaaac | aaggctaggt | 360 |
| agggttaacc | cttaggccaa | aacccacatc | gataactcta | cataaaagga | gacaatagat | 420 |
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| aacaaagtat | gggctttagt | tgaagcttca | aaggatataa | aaccaattgg | ttgtaaatga | 120 |
| gcttacaaga | aaaggattgg | agcaaattgg | aaggttgaaa | cctacaaagc | tcatcttgtt | 180 |
| gccaagggat | atgtcaaaag | taaggtatag | attatgacaa | aacttttctc | ccgtggcaat | 240 |
| gctcanatca | attcggattc | tttttgctat | agtagcatat | tatgatcatg | anatatgaaa | 300 |
| tggatatgga | aaatggtttt | acttaatggt | gagctaaaat | aatgtgtata | tgacacaacc | 360 |
| tganggatca | caccttgtct | g | | | | 381 |
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| tgcagggtta | attcaggatt | atggtagcgc | agaactattt | gaggtatgta | tgtaattctc | 60 |
| atagacttta | atttttatgc | ggtgtaaaaa | attttacatt | gtcaatcaat | cagaaatcat | 120 |
| catcagtgtg | atttttaaca | tagttattgt | aaaagtaaac | aaatttgtca | tattgtgatt | 180 |
| gaataacagt | gtaaagagtg | cataaactgt | ttactcattc | ttataatttt | acaactttat | 240 |

| | | | | • | | | |
|-------------------------------|---|------------|------------|------------|------------|-----|--|
| ctgcatgagc | taattatgta | gcttgaaatt | gggggcatga | gaactataag | atagtatact | 300 | |
| aggagtggat | ttataaattt | ctttgattta | aactataatt | ttagagtcat | tttgctatca | 360 | |
| agttgttgta | ctgcactgat | taacttatga | acttatggct | gcattccatt | gtttctgcaa | 420 | |
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| ttctttttac | atggagtctt | acacaagttg | caggaatgca | acggactcaa | cggtttctta | 60 | |
| actgcagtaa | ccaactactc | agtggcttaa | ctaaccgccc | ctaggttcat | acaaagtgtg | 120 | |
| acctaactat | acatgcatgc | agtcacgtgc | ctatacgtac | tcctccaggt | gccttggtct | 180 | |
| cgtgatactc | cttcatggtc | gtggctcgtg | tgttgcaaga | tgtggagctt | cattcatgtt | 240 | |
| gctgctgcta | acatcatcca | cccctgagaa | aaccaccttg | tgctcaagat | ggtgggcatt | 300 | |
| gcaaacatct | gtccatttct | cccatgtcga | gt | | | 332 | |
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| tgtgtcacga | ttcactgtga | cagtcaaagt | gtcattttct | tagaaaatca | ccaaatgtac | 60 | |
| catgagagga | caaagcacat | agatgtgaaa | ctacacttca | tcatagatgt | gattgaatct | 120 | |
| gagaaggtga | aggtggagaa | ggtttcaaca | aaagaaaacc | tggctgatat | gttcacaaag | 180 | |
| tccctctcta | gtgtcaagtt | caagcactgc | ctggacttga | tcaattttga | agatgcctaa | 240 | |
| agcagattgg | tagaagtgca | gccctaaatc | acaaggtaga | cacttgctga | tttggagtca | 300 | |
| aggtggagat | ttgtggtgtg | tgactcaaaa | tcacattggc | tcaagtgaga | aggttttaaa | 360 | |
| gtggtgttgt | cataactgtg | ttcagtcatt | ataattgaat | taggtttcac | accaatgtat | 420 | |
| agtc | | | | | | 424 | |
| | | | | | | | |

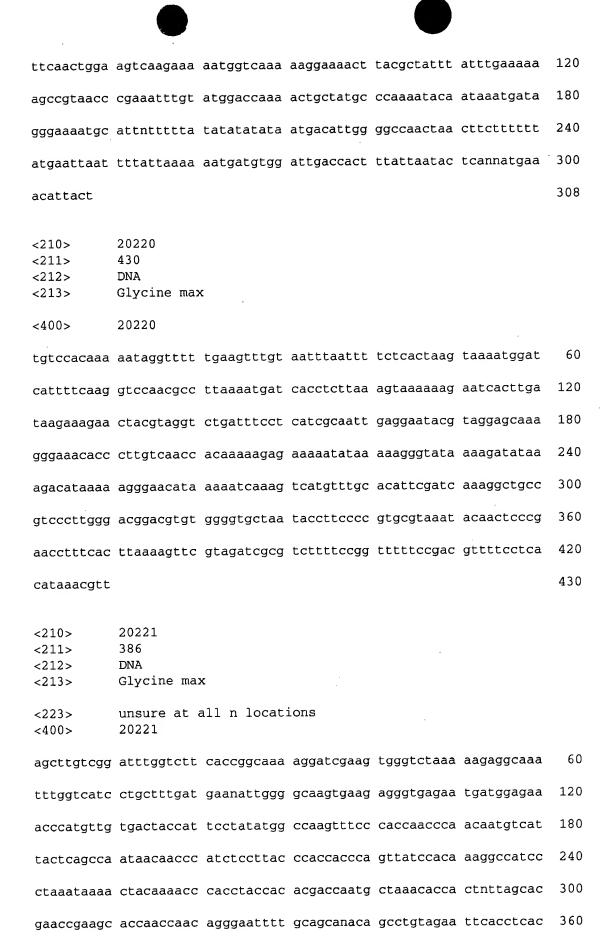
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| atctttatac | agtttatctt tcttacactt gagttttgga agaccaatta ctaagtcttt | 60 | | | | | | |
| cctaattaga | tgatttaaat gatgcatatt aatgtgtgca gtcctacaat gccacaacca | 120 | | | | | | |
| tgaatcatct | attttactca ccaagcaact tagttcatga aaaacatgct tgctcattca | 180 | | | | | | |
| acatatagat | gttacctatt ctcttaccaa tgtggataac ttaatcagat atggcttcac | 240 | | | | | | |
| ttataaggca | tcaatttcta ttaaattcaa tcttgaaatc tttatcacac agttgactaa | 300 | | | | | | |
| tgctaagaag | attatgctgt agtccatcca tatataacat attatntatc tngaatttgc | 360 | | | | | | |
| gttgattcct | tatatntgct tctcccattt atctctcttt ggtaattgct ac | 412 | | | | | | |
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| | taagaggett cetetagaag etteetegtg aettettega gaagetttet | 60 | | | | | | |
| | ctttgagaag ctacatcctt atctatccac ccctctatta actaaattaa | 120 | | | | | | |
| | aattattacg gatgaaaata acgcaacaaa taatcaaaca tcaaacataa | 180 | | | | | | |
| | tatatatata tatatata tatatata tatatata tataata | 240 | | | | | | |
| | cccacccttt tagaaatttc gtgctcaaaa tttaccttac tcaaacaagg | 300 | | | | | | |
| | ttctcgcatc tgactttcta attcccacgt ggcatcttct cctgatgcac | 360 | | | | | | |
| | caccitigace aacggaatet cittecetet taggigitggt glacgeetat | 420 | | | | | | |
| | | 430 | | | | | | |
| cctcgatcct | | 150 | | | | | | |
| <210> <211> <212> <213> | 20212 417 DNA Glycine max | | | | | | | |
| <223> <400> | unsure at all n locations 20212 | | | | | | | |



| ggacgatata | gattctttgt | ataccctttt | aagatcttca | tgtatcgctc | aaccgggtac | 300 |
|-------------------------|---------------------------------------|------------|------------|------------|------------|-----|
| atccaccgta | gataacagga | ccacaacatt | tgatttctct | gacagatgca | catcaagtga | 360 |
| atcatgat | | | | | | 368 |
| | 20215 420 DNA Glycine max unsure at a | | ions | | | |
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| aatttccctt | ggttatttgg | ctctccattg | atgtgttttg | gtgctttagt | tgctcatttt | 120 |
| ttgcaaaatt | cgtgaagcaa | tttgcatcta | aatccatgct | tgttttgtgg | agttgaggat | 180 |
| ttgaatgaga | aggccttagg | cctatgttgt | attctgaagc | aatggggcat | gccacattgc | 240 |
| ccccattctc | ttgcaattta | tgtccaaaca | tgtgcccatc | aagtgctcgg | tgaaatgccc | 300 |
| caatgatata | tgaatatgat | tttgcaaaat | tgggatggtg | gggctgtttt | gtgtatgtag | 360 |
| agacagcata | ggaaagtcga | aatagatgcc | caaatgcaat | cccaagctta | ggaacccaaa | 420 |
| <210> <211> <212> <213> | 20216 327 DNA Glycine max | × | | | | |
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| actgcagttc | tgatgggggt | aggaaaattg | atgaacaaaa | tgaattacaa | acaaatattc | 120 |
| aagaaaaaaa | aaaggatagc | acgtacagat | tagtttatta | gacattgtaa | aacaaaagct | 180 |
| ataaaagaaa | acactttaag | tatcaaaaag | ggacacatta | agtatcaaaa | agggacaaag | 240 |
| atgaaggaaa | cacttaggtt | ttaaaacttt | gaagttgttt | gtagcctaca | aaaatgcatt | 300 |
| gatttgataa | gagtgttcaa | gttatta | | | | 327 |
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| <213> | Glycine max | |
|----------------|--|-----|
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| aacctgtaaa | aaaataaagt gtttggtaaa gaaaaaaaa tgggtatttc atttcatacc | 120 |
| acatttacac | aatgctacat tgtaacaagt cttacaggaa tttttcagc aagaagcttt | 180 |
| cgtattccat | gagetetaga etecacett acatecacte etgaaacata tteaggataa | 240 |
| agcaccagaa | gctccatgat atttgtaaga tacggattta aaaatccacc aagcttgtcc | 300 |
| acaactgcct | ccaaagtgat gagaacataa aaatgggatt catttgaggc tgataaaaca | 360 |
| tcaatggttt | cgggtttcat atcagccaat actcgacgag atgacttcat cacattgtcc | 420 |
| ataatcttgg | g · | 431 |
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| gtatgtcaat | atgtccaccg atgaaccett ggaatgagac accatcaagg gccgtaggga | 180 |
| acaacacgac | ctaagcgaat tttcgggggg ctttatcagg caacaatagt gagctcaagc | 240 |
| tcccaagagg | tgaaaggaat catcacgggt caaaggcatg atcttgatcg acgagctgca | 300 |
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| gt | | 362 |
| | | |
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| <210> <211> | 308 | |
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| <400> | 20219 | |
| tttttttatt | tggcttgatg aattcttaaa agaaattaat tgaaaccaat gacgattaac | 60 |



<5772>

<510>

₫35 5055₫

| | | | | | , , , , | |
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| | · | • | | | | |
| 198 | | | | | | ב |
| 360 | аатаатааат | агагдааааа | cancatttat | αςσεεεαεας | гаасаааатт | тсаааататт |
| 300 | ttctaattgg | agtttgtttc | dcrccdsccd | αςαααςςςος | atgtagaaat | tecttatage |
| 240 | аасадаадса | гсггадссва | aatgaaagtc | адасдсаваа | aactaacgga | ааттаасааа |
| 180 | ассдасдсед | сғаағаяааа | ttatctaatt | ссгататааа | тсадттата | cattgactat |
| 150 | tacatgttga | тстававдст | atatatgatt | ರಿದಿಂತವಿರಿ | гагдагадсд | ссаастдаса |
| 09 | ttttaagaat | агаагяагяг | ссаааасаса | aggacataga | cactgatcat | agcttttatc |
| | | | suo | ill n locati | SOSS3 nuente et e | <222> <000> |
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| 360 | crødrddard | ταςτταττςς | cttcttccat | адсавадссс | tcatcctggt | ratccaaggc |
| 300 | ccsagcttct | gggcgccctt | τοςτοσττα | כלכלללכלל | οτττοτος | cccccccccc |
| 240 | τττοαταοτο | attectteaa | σοσστοτττ | tetaetete | acttcaaagt | tatgagacac |
| 180 | атдааадтст | actttgatga | acttgttgta | catgtgtaac | гааатадаад | асстадіста |
| JSO | сваддаадст | дваадсссс | קבבבכבפפ | tctcaaggaa | ರವನಿತನಿಗಳ | дээдээдбееб |
| 09 | dcffcfcaag | сгаадсссга | ααςαςαςς | ttatgagagg | dragacread | гсгсээддэд |
| | | | suo | | 20222 unsure at a | <223> |
| | | | | | дуусіпе тах Сіусіпе | |
| | | | | | ANG | |
| | | | • . | | 431 | <5112> |

20222

<57.0>

| 150 | שכשבשבבמבב | ಕರತಿಕರತಿಕರ | agacagctta | ಕಕಿರ್ದಿಗೆ | агасадсаад | acatatatac |
|---------|------------|------------|------------|--------------|----------------------|------------|
| | | ctttattaca | | | | |
| 09 | Deeteesti | 6061161110 | 1010104001 | 1200112000 | 040000000 | |
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| 240 | tttacatgtt | gattctggat | στατταττ | αςσετταετα | cdattaaacc | cddadttata |
| 180 | catcctgttg | аддявасавв | гадагааааа | cdrcddrdsc | ctttaaatc | гяддггггд |
| 150 | ааастутдаа | rrdcrdrddd | дгадггсгга | בבקבבפבבב | cdaattctga | ddgrcrdrrr |
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| 432 | | | | | O.D | 0044000420 |
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| 007 | · | | | | | |
| 98 | ccaccactgc | ೦೦೪೦೪೦೩೩ | дгааатуаст | дасааатсаа | ೧ ೧೮ ಇ೦ ೧೮ ೧೮ | ggaatcatat |
| 300 | ccccdcagaa | вадавдсвва | cdcsdsfddc | grgacacaac | ccttgaaacc | аадааасаса |
| 5₫0 | αττττατοσε | сгдгагдсаа | acctaaagaa | caccaattgc | crasssesa | ссааттстаа |
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| | | | | | Glycine max | <213> |
| | | | | • | ANG Yem Agiby[9 | |

| 360 | свавасваду | ddggcdccdg | dscccddsss | sacagagacc | ಶತತಂಡಿದೆದಂಡಿದೆ | ದಿರಿತಕಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕ |
|------|-------------------|----------------|--|-------------------|----------------------|---|
| 300 | ಆ೦೦ಡಿಆ೦ಆ | cccccddcs | ccsgagggss | acctcgagaa | ccssdddsdd | ಶငငငင့ಆಧನಿದ |
| 240 | cdødddcdøc | ಶತತತದತರತರ | cccddddsss | ತತತ್ವತ್ತರಕ್ಕ | cdcsssagss | csgagacccc |
| 180 | ссясствата | ದಿನಿತಡಿದಿದ್ದರು | ಇದಿ cಡಿಡಿcಡಿಡಿಡಿ | cccscdssac | actaatttcc | ತಂರ್ರತಿಕಾರಿತ |
| 150 | яяясяддд <u>а</u> | ccdcdcddsc | ಇ ೧೮೮೮೮೮೮೮ | agaancgcgc | tttgaactag | ccscddsdcr |
| 09 | ದರವವರರರದ ದರವನ | agacctcaaa | ಕಡಕಿರ್ಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿಕಿ | ಕರ್ಶಕ್ಷಚಾಗಿಕ | ಶಂದಶತದೆಂದಿ | ddgccdgfdc |
| | | | suo | ill n locati | SOSS8 nusnie st s | <223> |
| | | | | 3 | Glycine max | <213> |
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| | | | | | 20228 | <210> |
| 9₹€ | | дясярд | двагдагдап | ಕಡಿತ್ತುತ್ತುತ್ತ | aatccttctc | attgaggaca |
| 300 | ddadaatccd | gatgcagaat | ατατατυτ | tetetgattt | асстссвату | tgttagttcc |
| 5,₫0 | агдадгагаа | dødcrdcccd | грасвавдър | atgacaacgc | дааадааたса | tcaagtgctt |
| 180 | atggaccatt | ccssddddsd | aaagcttcaa | ತಕಕರ್ರತಿಕರಿ | rttccggaac | ಕಡಿಕಡಿಕಡಿಕಡಿ |
| ISO | tgcacatgag | εαααεεεααα | ತರ್ದಾಗಿತ್ತು | ттдтстсаа | ತರ್ವತಿಕಿತ್ತರ | свасвавдду |
| 09 | ctaaacaagc | аааадссасд | ರತಿತತಿತಿತಿಕ | atcaaattga | ададгсааад | атсттагдад |
| | | | SIIO | י דו זו דחממרד | SOSS1 nuante et e | |
| | | | 540 | | | |
| | | | | | DNA Glycine max | |
| | | | | | 978 | |
| | | | | | 20227 | <210> |
| ££₹ | | | | | 282 | gcctgattat |
| 420 | grgaatagca | ttgaagcaat | аадастддат | ccrcrrardd | ραςαςςαας | тсаатстста |
| 360 | ಶತತಿರ್ಭರತನ | гсгсгсддуг | двсядгядсг | aatttettg | ತರ್ರರತರಿಕರ | aaattaatga |
| 300 | ctgatcatta | гатсадстдд | ггадгсаааа | ccttaaggac | гсггяддссг | атааастдаа |
| 240 | cffddgdcgf | агдсяссяяд | tataaatcgt | адѣѣдаадса | ctccccctca | aattctaaca |

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| ccacata | | | | | | 457 |
|----------------|-------------|-------------|------------|---------------|------------|-------|
| attatataaa | τασοτατοστ | tgatccattg | ςδαδαςδες | тасатадсса | rdcrrcsddc | ₹5(|
| satatgtaca | gagaactaca | atcaattaga | аатааттсст | cscsttttgt | cctctccttt | 390 |
| асавстсстд | tctattatgg | acatctattt | ccsctgttgt | accttagact | aagcgaaatc | 300 |
| ctgtactatt | ttatgatagt | дэдгээгдгд | rddsscardc | адаасаддса | ggaatcaatc | S ₫ C |
| gatacctggg | версессе | аттстдтатд | атстсстват | agataattat | gragttaatg | J8T |
| tgacaaaga | aattggaaga | аатттатаа | аддагсггаг | адгдзэгдэг | астдстдааа | 150 |
| застсадстт | csdtggagag | agatctcatt | аггсгээссс | ttgtattagt | draagggaga | 09 |
| <007> | 20230 | | | | | |
| .CT. | | _ | | | , ė | |
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| | ANG | | | | | |
| | 724 | | | | | |
| <210> | 20230 | | | | | |
| ссдссвдсс | аддсядряяд | £daagttgag | aaagatatca | сгаст | | 50₹ |
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| rtddrdgcrc | tagcataatc | аассдаддас | catatttct | ರ್ಡಿತ್ರದ್ಧಕ್ಷ | acattgctca | 300 |
| адссттава | аватавсссс | attattacat | гдссгээдсс | агдадагага | cataagtgga | 5₹0 |
| eagcagtat. | дсяяддсяся | ааддаассда | gatttctcca | сдссвааста | аататсаддд | 180 |
| ಕ್ಷಕ್ತಿತ್ತುಕ್ಷ | taggacaccc | aatatccaag | тдатстсват | тсасаадста | כבבבבבבב | TSO |
| ומכבבבפב | ataaatgacc | aataattata | acagaatttt | atctttcaa | tatcatttga | 09 |
| . <00ħ | 20229 | | | | | |
| | unsure at a | ll n locati | suo | | | |
| <513> | суусіпе тах | | | | | |
| | ANG | · | | | | |
| | 907 402 | | | | | |
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| -010 | 20229 | | | • | | |

16⊅ ವಿಡಿತಕಾರ್ಕಿಂದ ಅಭಿಕಾರಿಕಿತ್ತರೆ ಕಿಂಗ್ ಕ್ಷಣಕ್ಕೆ ಕಾಲ್ಲಿ ಕ್ಷಣಕ್ಷಣಗಳ ಕ್ಟಣಗಳ ಕ್ಷಣಗಳ ಕ್ಷಣಗಳ ಕ್ಷಣಕ್ಷಣಗಳ ಕ್ಷಣಕ್ಷಣಗಳ ಕ್ಷಣಕ್ಷಣಗಳ ಕ್ಷಣಕ್ಷಣಗಳ ಕ್ಷಣಗಳ ಕ್ಷಣಕ್ಷಣಗಳ ಕ್ಷಣಗಳ ಕ್ಷಣಕ್ಷಣಗಳ ಕ್ಷಣಕ್ಷಣಗಳ ಕ್ಷಣಗಳ ಕ್ಷಣಕ್ಷಣಗಳ ಕ್ಷಣಕ್ಷಣಗಳ ಕ್ಷಣಗಳ ಕ್ಷಣಕ್ಷಣಗಳ ಕ್ಷಣಕ್ಷಣಗಳ ಕ್ಷಣಗಳ ಕ್ಷಣಗಳ ಕ್ಷಣಗಳ ಕ್ಷಣಗಳ ಕ್ಷಣಗಳ ಕ್ಷಣಗಳ ಕ 08₺ ರ್ಥಿದಿರಿದಿರಿದಿರಿತ ರಡಿತತತತಡಿಗಿ ಕಿರ್ದಿಂದರಂದರ ಕಿತ್ತಾರತಕೊಡ್ಡ ಕಿರ್ದಿಕಿಸಲಾತ ಕರ್ನಾರಕಿಗಳು 450

| 09 | сстттаат | ctactaacac | аасдііі | atgagaagct | agcagcttct | gacttcttt. |
|--------------------------|--|---|---|--|---|--|
| | | | | | 20233 | <004> |
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| 300 | ссясяядсся | ggatgaaatt | tcattgaact | accaattaca | tacagttgtc | tatttgacat |
| 5₫0 | ತಿತಿರ್ಧಿ ಕ್ಷಣಕ್ಷ | ссссадсаа | tacaaaatta | даддсасата | פּמרבפרפרפ | свавдавдаг |
| 180 | gagcatagaa | tggaaagatc | aatagtatta | tcaaactttt | дсгээээдэг | асвадссвад |
| 150 | tgttatcatc | даттаасааа | csataggaga | асдсаасааа | aggtatctcc | ದಿಡಿತಡಿತತಿತಿತಿ |
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| | | | | | _ | |
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| | | | | , | DNA | <212> |
| | | 6266222 | ananannaan | | 20232 417 . | <210> <211> |
| 80ħ | | | | peonttatet | DNY 50535 ££dcccsdff | c212> <211> <210> |
| 80ħ 09£ | canaagacag | cactagtctg | | peonttatet | DNY 50535 ££dcccsdff | c212> <211> <210> |
| | α, | | tacaatgatc | taaggcacca peonlistei | DNY 50535 rfdcccsdrr | <pre><210> <211> <210> <310> </pre> |
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| 360 300 340 | ассеседенс сасседддаа асгдсадаас | acaccagttt ctaagaagtc | аддсгссвас сваагтгдгс дддаваггас гасватдвгс | дстдтасаа сстававаат стдасаваад тааддсасса тататтпсад | DNA 50232 cltcctccadt cagagccadt cagagccadt | caggaaaatt caatcagctc ggaatagca tanatgccat caggaccaca |
| 360 360 340 380 | agcaccaata actgcagaac actgcagaac | аадагаааст асассадггг сгаадаадгс сасгадгед | туссааатас дуустссаас сааатттутс ууудааттас тасаатуатс | ccaattcctg gctgttacaa cctaaaaaa ctgacaaaag taaggcacca taaggcacca taaggcacca | DNY 50535 cadadccadr cadadccadc | gagaaaaga caatcagctc ggaatagca aaggccaaca tanatgccaca |
| 300 300 540 180 | agcaccaata actgcagaac actgcagaac | аадсссса аадагаасс асассадст сгаадаадс сассадст | туссааатас дуустссаас сааатттутс ууудааттас тасаатуатс | ccaattcctg gctgttacaa cctaaaaaa ctgacaaaag taaggcacca taaggcacca taaggcacca | DNY 50535 cadadccadr cadadccadc | gagaaaaga caatcagctc ggaatagca aaggccaaca tanatgccaca |
| 300 300 540 180 | agcaccaata actgcagaac actgcagaac | аадсссса аадагаасс асассадст сгаадаадс сассадст | сатуссваса дудаватта дудаватта саватттут саваттут саваттут саваття саваття саваття саваття саватия оз | ccaattcctg gctgttacaa cctaaaaaa ctgacaaaag taaggcacca taaggcacca taaggcacca | DNY 50737 cagagccagt cagagccagt cattcttctc aggagctagt aggagctagt aggagctagt | <pre><400> <400> <pre>cadgaaadt cadgaaatt cadgaaatt daaatagca aaggccaaca aaggccaaca </pre></pre> |
| 300 300 540 180 | agcaccaata actgcagaac actgcagaac | аадсссса аадагаасс асассадст сгаадаадс сассадст | сатуссваса дудаватта дудаватта саватттут саваттут саваттут саваття саваття саваття саваття саватия оз | agaaaaagag ccaattcctg gctgttacaa cctaaaaaat ctgacaaaad taaggcacca taaggcacca | DNA 50232 417 cagagccagt cagagccagt cagagccagt agataccagt | <223> <400> alctataagt caggaaaatt caggaaatt caggaatt caggaaatt caggaatt caggaaatt caggaatt caggaaatt caggaaatt caggaatt caggaat |
| 300 300 540 180 | agcaccaata actgcagaac actgcagaac | аадсссса аадагаасс асассадст сгаадаадс сассадст | сатуссваса дудаватта дудаватта саватттут саваттут саваттут саваття саваття саваття саваття саватия оз | agaaaaagag ccaattcctg gctgttacaa cctaaaaaat ctgacaaaad taaggcacca taaggcacca | DNY 50737 cagagccagt cagagccagt cattcttctc aggagctagt aggagctagt aggagctagt | <213> <213> <213> <223> <223> <223> <223> <223> <223> <223> <223> <223> <223> <223> <223> <223> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233< <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> <233> |
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| 300 300 540 180 | agcaccaata actgcagaac actgcagaac | аадсссса аадагаасс асассадст сгаадаадс сассадст | сатуссваса дудаватта дудаватта саватттут саваттут саваттут саваття саваття саваття саваття саватия оз | agaaaaagag ccaattcctg gctgttacaa cctaaaaaat ctgacaaaad taaggcacca taaggcacca | DNA | <pre><212> <212> <212> <213> dagaacaaca dagaacaaca caatcagctc dagaacaaca caatcagctc dagaacaaca caatcaaca caatcaacaaca caatcaa</pre> |

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| aaaccatacc | ςοερετεααα | attctatacc | гсгггядсгя | caatttccaa | дагадгосда | 240 |
| caattatcac | ccscddrdds | ададссдаа | дваатсстдт | αστασοσεςς | ದಿದಿತಡಿದಿತತತದಿ | 180 |
| attgtctccc | ссдсссвата | грагдассад | ccdrtdaagt | derreseerr | rddddacrrc | 150 |
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| гддясгггя | драгадгдг | сдддрдачдч | ддясягдггя | αρασαρεασο | cffccfccdc | 300 |
| gattcatgcc | crdudadrdd | tecetteaae | gcttcaccag | аааттдаадт | ್ದರ್ವತಿಗೆ | 240 |
| ςαςααςςς | свадсаасдд | atataaatcc | tgcctatagc | τασσεσεταα | двадассдта | 180 |
| гсдагадавв | агаагааааа | aaatcgacat | recearteag | ataggccccc | acacttgcaa | 150 |
| тссттта | gegteaeatt | гваадссдад | crcdattttg | drdcdsdccr | ttgatggtac | 09 |
| | 20234 nusnxe at a | ill n locati | suo | | | |
| | Glycine max | 3 | | | | |
| | 403 ANA | | | | | |
| | 20234 | | | | | |
| садъдсасаа | aatttcacag | ರಕರವಿರವಿರ | | | | 388 |
| aattcaaagc | ctatttaag | τοτατοττατ | gcaaattan | ರಿರ್ಧತರತಚರ್ಗ | gtataataac | 360 |
| сгаадстсаа | atccactcac | rcdcdcrggd | cdcdaddafd | дсдсгаадсд | ರತಿಕಂತಿಕಂಡ | 300 |
| дсдядярсяд | гдгдсгээдг | дсядгядггд | тсттсявсся | ddcrcgdcgc | асдастадтд | 240 |
| acatcaagca | тааттастаа | атататата | асасасасас | ассадддсдс | гасастаадс | 180 |
| аастааассс | acctccttga | aaataattac | ggataaaat | аасасаасаа | ататаатсаа | 150 |

| | | | suoj | all n locati | 50538 nuante et 9 | <2223> |
|-------------|------------|------------|------------|--------------|-----------------------------|-------------------------|
| | | | | > | DNA S0238 | <210> <211> <213> |
| 373 | • | | | | £ад | aatactgttc |
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| 300 | atatattatt | caattgatca | ttaaattota | aagttattt | tcttaaataa | ttgtttaatc |
| 2₫0 | agaaatatgg | בבבבבפקבבכ | tttaaatata | taattaaaca | сатасаааат | ctagtattat |
| 180 | gccattagaa | αρροστρορο | ttgttccata | tccacggacc | аассасадсь | cdcstdcctc |
| 120 | agtacatccg | ттдсятсявя | агавссггав | адссяяссар | гаарссяваа | tttatggaat |
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| 300 | agtaaagega | сссстадсаа | ааааааасас | ататаатаат | taaaataata | รรรรรษ |
| 240 | дггддзээгс | ccdsaaccac | ερεαθεσαεα | attccgaccg | ассававсав | дссаассссд |
| 180 | tgtatcatct | rcdrdrddgr | tttccaccga | taaaataaat | ааастааааа | ττιςςcdctτ |
| 150 | atttaagtca | гаадссаггг | trgacgtgct | ataccccctt | ttcactttt | acacaggtca |
| 09 | קדמבלבמב | tttgctacat | cddgccffgf | дгасссдсдд | тетоттеотт | сдсддггсва |
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| 240 | tctttagaca | дваватддгг | ававтттса | аааатсдтда | gaaatttta | aacttttgga |
| 180 | аааасдасда | адсагдгагд | атдгтгдааа | τταατοττασ | acaaaattga | гатдсттдаа |
| 150 | cattactttc | cddcdfcgfd | τασταττατ | ггдгэдссдд | аааддтаата | Laaaaaggga |
| 09 | ттатсаваса | ttcccttatg | сттататсаа | dddcddaact | catgcgaaga | равсвавадд |
| | | | suo | ill n locati | SOS∢O nuenxe a¢ s | <223> |
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| 282 | | at | dødrrdrosr | τοτταττατ | ςτςττάτττ | τατταοττατ |
| 240 | αδρορραφία | בפפרנתרננ | адсгигаасс | ccattggttt | сдссававсс | аааассдссс |
| 180 | taatccttga | аассдссдаа | tcattgcttt | tatgtccttt | agcttgcttc | tttaagttct |
| 150 | taagtttctc | agagccattc | астстастст | 1111111111 | τττυτττασ | rrdrrdsdrc |
| 09 | ತರ್ನಿತಡೆತನ್ನ | caagttccaa | аасдсдаааа | attgaaaatt | ctaaatttca | מבכבבמבבבכ |
| | | | suo | ill n locati | 20239 nnsure at s | <223> <400> |
| | | | | 3 | 20239 DWA Glycine max | <211> <211> <213> |
| 4 29 | | | | | | псдааасаа |
| 075 | าาทาาททาทท | annnaannn | ກກຄວວກວກວ່າ | מרממרומחרמ | ממארממממממ | רממממחמרמר |

| <004> | 20243 | | | | | |
|-------------|------------------------------------|-------------|------------|--|----------------|-----|
| <211> | 414 DNA Glycine mas | > | | | | |
| | | | | | | 898 |
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| gacttaatcg | מכפרפרמרפר | садаваядстс | tttcatgaac | tgattagcac | atgaatttt | 300 |
| caattaagtt | тсаадастса | agttccatga | atcaatatca | agattcaaga | атсаададаа | 240 |
| grtcaagaatt | дватсатсва | cactccatgg | ctcacatgga | aatttgattt | свададссва | 180 |
| ctctaaagtc | atgagcactt | catgatacaa | acgatgatga | tctcacgact | caatacatga | TSO |
| tcatgatgaa | ссаадастда | ttcagagagt | ttcgatgttt | ್ ರತ್ತತ್ವಾಗ್ಗೆ ರತ್ತತ್ವಾಗ್ಗೆ | £dacaaaaag | 09 |
| <004> | 20242 | | | | | |
| <511> | 20242 DNA Glycine max | > | | | | |
| 999 | | | | | | 243 |
| ರಿತರಿತರಿತತರ | гдссгээддд | gctgacccct | τοςττοττας | οττοοτοσος | tatttatagc | 540 |
| rcddssdcdc | сғсддсғғв | attttcttca | cggaaacaat | rtttccaagc | caattcgaaa | 180 |
| acgccgaaga | асддссдвая | гсгггдсдээ | attecteace | даааасусса | ೦೦ದಿತಿತಿಎಂದರ್ಗ | 150 |
| cagettttt | cdøddføcff | асссддгдаа | дассдаадаа | ರಡಿಕಿರಡಿಡಿಡಡಿಡಿಡಿ | cgaatgaaga | 09 |
| <004> | ZOZ⊄T | | | | | |
| <211> | 20241 243 DWA Glycine max | 3 | | | | |
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| agaaacatta | саадаасаէէ | nttattggcc | tcatatgana | ತನಿರುತ್ತದ್ದರ | agaaantttg | 420 |
| מכבכמבבנמב | даадсасатд | aaggaggttt | аасудуусас | ς ε ε ε ε ε ε ε ε ε ε ε ε ε ε ε ε ε ε ε | аааадастст | 098 |

agrictigicas ngalgigagic teaangalgas dacgggcgtig tglaactaaa cictaactaa 540 agrictigida gacatactic anagitigica tictatecet cicaacaaag citcicaagig 180 agrictigicas agaagcatic gacagactic cicaacaaag citcicaagig 180 agrictigicas actatacat agaagcatic gacagactic cicaacaaag citcicaagig 180 agrictigicas ngalgigagic teaangalgas gacgggcgtig tglaactiga cictaactia 540

<213> Glycine max

<210> 20245
<211> 291

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ntraaattea aatttggaaa ttgcaaattg tttcagattt caatttagee actggtaatt

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<223> unsure at all n locations
<400>

<213> Glycine max

<211> 417 <212>

<210> 20244

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| | Xom anicy(2) | <£15> |
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| | ANG | <515> |
| | 917 | <511> |
| | 20248 | <510> |
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| | | | | | • | |
|----------------|-----------------------------|--------------|------------|-----------------|----------------|-------|
| catta | | | | | | 392 |
| гвасдаасдс | rdddcgrrcd | rrrracagag | deagetteat | αςααςςςςς | cragaagctt | 3 6 0 |
| tcatatcatg | дасяядгагс | ttgtggagtt | dcsdsccsns | tttcgagaag | atccaacggt | 300 |
| сядяядсьс | attaagaggc | ttatagcaca | grccagacar | сггсгаааад | atcccaacgg | 540 |
| cgttttacc | дяддсядсгг | cstdtsgctt | гсгсгядяяд | cttcattaag | aggetteete | 180 |
| агсггагава | dffdcgggcc | aaatttagag | аадатссаас | ggttaacgaa | ರಿರ್ಧಧಿವಿಧಿತತದ | 150 |
| адстсстатс | acacttcaga | aatcttctca | аадатсссаа | ಂಡರ್ಥಿಂತರಿತ್ರಾಂ | atggaaagt | 09 |
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| aatgtatatt | аасдссдааа | дгравсагсд | дгссгасгд | аваассдагд | тгдггагсаа | 360 |
| aaaattacca | асатсддтта | татааатаас | cdstdttdct | аасасдаасс | эсясссядяс | 300 |
| aagtaccgac | дггдагадга | ttatcgttaa | cstcggcttt | £dagaaaccg | атдтгаасдт | 240 |
| aatatgtttt | ctacatcggt | tatttatgac | tttcaacatc | ddcffffcag | ccdatgttga | 180 |
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| αςεεαεςςαα | двасадттат | gcatgagttc | atagatgtgt | ασεεεσεεασ | drccrrccds | 09 |
| | 20246 unsure at a | ıll n locati | suo | | | |
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<007>

20248

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<210> 20253

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| 240 | catctataat | ಕ್ಷಂತಿತ್ತುಕ್ಷಣ | свадтатуда | gacaatgcaa | αρρααρρααρ | ttcccttgga |
| 180 | gacacttccc | асдссдадда | carastraca | аааадстдта | cactatcgaa | τεοοστάτατ |
| 150 | acattcggtt | cfdcsaggat | ctttctgaat | ddccfacaga | гсддэдсээс | rradssssa |
| 09 | rrrggddrd | actacggccc | αςςςςςςς | гсстааатса | ctcacttcta | ಶರ್ಧದಿತಡಿದಿದಿದಿ |

| 3Jycine max | S13> |
|-------------|-------|
| ANC | :515> |
| 158 | :S11> |
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| | | | | 12505 | <001 |
|-----------|---|-----|------|---------|-------|
| locations | u | all | at | əznsun | < 522 |
| | | x | em € | суусіле | <£12 |
| | | | | ANG | <212 |
| | | | | 412 | <112 |

20251

<510>

atatgetase tigateceat atetaceat taatecata gratecata eccetage 414 assacease aggaatitig eageasaag cetgeaggat teaceeasa teegatgee 360 cettaatee accaeasa eccetageg coettageg 300 attateceas eageatee ectetecet accaeacac cagtratees 240

<515>

<511>

358 DNA

catttgcatt acaacattca tggccttgca ggagaatacc cgcgcaaaca tttgaaggaa agctigiges iccigiacce igaigaggat gacgeataig itcitataac iggacicate 09 20255 <007> Glycine max <213> <517> DNA 797 <5112> <570> 20255 Ţ 451 ggcaaccttg atatcaaccc ttatgaataa cattgattga ccgactcttt tttttatata **₫**50 gcatcettaa ttaatcactt gatataatta atatgatgtt gaaagttaa teetgetega 360 वर्ष्वरेवववदवट रंबटर्रपुर्वतेव वर्राट्यववटव ट्यूप्ववववदव रंप्यववर्ष्व वर्ष्यववयदेव 300 aaccttaatt tacgactagg ttgcgtttca cagaattagt agtatcttgt ttaattctac 240 tettggatge tagactgatt eteatatgag ttaattactg teactaatte caactettea 180 ttttggggtgg ctctcacgga tcaatgtcta cttaccaagt caaatacatt ttacccttcc 150 taacgtgtag tegtgaaata tgtaaaaat tecatttagt gactntetet agaetecaae 09 7025₫ <000> unsure at all n locations <222> Glycine max <213> <515> ANG <511> **451** <570> 2025₫ trectitic tittraatat tgatgeatit cttatagitt ataggaagta agtgtaat 328 agcttaacaa ataacaaaga tggagcacat tttaattatt ctttataata atggattcct 300 decocesate atgattasas aasagtacat gotacgitga gitaagagta ticcagaaca 5**₫**Q gaactataaa aaatacacgg aggaccaaca attcattgaa accataaaaa tctaagctca 180 टांबर्टाबर्प्याचे प्रवित्रवित्येव वर्तावेबर्प्यवेवर्षे एप्यावेबर्याचे प्रविवेववेबरेट 150 agcittatgg caattigatg gotacatgag cacttaacac acaccactta goagcoaaga 09 20253 <001> Glycine max <512>

<007>

unsure at all n locations <222> Glycine max <213> <515> **₫**30 <211> 20258 <510> gcacttcac 370 360 ddcffffatg gctactacaa ggtfgfataa tctagtatac ttcctatatg tgaaaataaa refecence earcreese areaatitee eteceteegg teacetaase etaacaceet 300 catgacctit tittatctaa gcaaattatt atatttattt ctcctaatga tatagagatt 240 tetectiett ecataceat atglecteet cacteagaat caaattaata ettetaagt 180 tergreeatt acacaaaat aacacatet aagcaactt aactgagtag agactagtte 150 градстрдся састсаваду дасатстсас стататстт агдававаст агсатутаса 09 <007> 20257 Glycine max <512> <515> ANG 310 <5112> 20257 <570> drdaatgtee taagaaacta tactgetett cagggtetet teattgatgg ga 232 dacatotaca gactadigg ttaaggaacg cgacaggata caacttatt tcacgaacct 180 वेषटट्वेषट्वे इम्ट्रिकेवेषेट व्वेट्वेवेवेषेवेष्ठे प्रवामानिक प्रतिवेषवेष्ठ व्वेट्वेष्ठेष्ठ व्येट्वेष्ठेष्ठ व्य 150 हतीबर्दाहर वार्षे वार वार्षे वार वार्षे वार्ये वार 09 20256 <001> Glycine max <513> <517> ANG <5112> 232 20256 <570> atgaccdtca ccagctgcga tgac 797 высустубень сестемень возыция возысовый возысовый сеторогы сеторогы высусты в rereatity tyageteeae catgaageee eeacatgtee atgatgatea catatatety 180

| 5₫0 | tgatactttg | ttccttttaa | даатстаата | cttcccdat | atattgaagt | tccagcagtc |
|-----|-------------|------------|------------|--------------|--------------------------|----------------------------------|
| 180 | астдсдссва | ttgcttcaaa | аатадатдса | адагадгсад | ddfdcffsds | ραδαρασρα |
| 150 | ರವಿಥಡಿತಡಿತಿ | аааатаатаа | catcatgcag | дгвасдадад | гадггссгда | tgaagacttc |
| 09 | acatttagcc | аадтусада | ctctatctga | agtttctcaa . | адвваварс | игдэдгсгээ |
| | | | suo | all n locati | SOSEO nuenze er e | <2223> |
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| 300 | cfffcfffgc | ссаатадаат | ಶತರತತಿರಿದಕ | aattattcta | tttgatgagg | caaattgctt |
| 240 | cagccagaac | tgttcaccaa | בנקבנפנקנ | дгггадаагд | agtacccact | эсгразарсг |
| 180 | gataagagca | tgacacatat | atgcctatgt | рсвсвавссв | грсрсягард | эссваваттс |
| 150 | gaaccttacc | ttatccatga | аддадсаггд | сваасттата | cctataccga | ttgtacagaa |
| 09 | catttttat | catcattatg | дгссгаасаа | стсаастааа | rdrdssdssc rdrdssdssc | адстттдаа |
| | | | suo | all n locati | SOS28 nuanke at s | <223> |
| | | | | 2 | Glycine max | <213> |
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| | | | | | 20259 | <210> |
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| 420 | gattntttn | tactatatgt | аассаассад | cscctttgtc | адгсаадага | attgttaac |
| 98 | стстдатсаа | гаадаагдад | сттаўааста | ττττοτταττ | ddscrcdrsd | дсяяягддя |
| 300 | agtgtttgg | εςςααςςςας | дгдагаасаа | atgagttacc | аасдассада | дссвадсьдс |
| 240 | ςςαςααατες | ttctcacaat | tecttegaae | ataatgactc | tttgcacaat | адагаттсат |
| 180 | ггядсгадга | tacaattact | ragagtttgg | בבבבבבכב | ггдаатдсаа | cacttgttct |

taccatcaat gacaatgaag tattgtgtgt gggatgtgt ttgactgtaa atatacttgg $120\,$

09

ratidotidg asaccascae tagataagat actotytin tigaatitat ctcacaagaa

<210>

| 907 | | tagaac | аадстасста | гсяагдссгд | сгадастатс | rrccrccagc |
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| 300 | aatgaagttt | дгсггддзэс | ддсгсагсаа | тдстттавая | гдгсггэгээ | дассассасс |
| 5₫0 | гадгггдсаа | crddrdsccc | drrdadacad | ಶငсಶರಿತದಿತರಿ | aaccagatcc | cctgcattac |
| 180 | ತರತ್ರತ್ತಕ್ಷ | ತರ್ಧೆದಿದಿತತದಿತ | tttgttccac | taagcaatat | дггдгссгээ | cctagccatc |
| 150 | csagtttcc | cttgtgatga | ctatgcttta | ಕರಿತಕಾರಿದಿದಿದ್ದ | acttgagaag | gcagaaatgc |
| 09 | аасдасассд | атдттаадаа | atatttatca | ತರದತರದಡಿದ್ದ | attacccaca | tacccgcact |
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| | | | suo | ill n locati | unsure at s | <222> |
| | | | | 3 | Glycine max | <513> |
| | | | | | ANG | <212> |
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| 240 | ггдсгассаа | дсссддааас | тагдастаст | тадтатарат та | cataagtttg | тдгаадгааа |
| | | сагддггсга | | | | |
| 180 | | | | | | |
| 150 | attcatatta | ctacacattt | ataacaataa | aaaatcccaa | ttctcttaaa | ttctattta |
| 09 | tttcggcatt | trccatatat | ттсяясттт | reatttacgt | дггсгэдгсэ | гадсттгдаа |
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| | | | | • | 352 | |
| | | | | | 20261 | <210> |
| | | | | | | rasagc |
| 925 | | | | | | ODEEET |
| 02ħ | ccdccsdcdc | caacggcaaa | ааасддсгса | гсападагса | датстатаас | асатдаааса |

acatgasaca atactetag aggrasat tettageac tacaggras greatect 300

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| 150 | ααςαοςστρα | сагсядагад | gccffddfdg | ದಿಡಿದ್ದೇಂತಡಿಡಿತಡಿ | rardrododd | гассгддада |
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| TSO | ttatgtttta | сатсававьс | татдтсаата | ςτεςτααττι | aaattttgg | ааасаатсса |
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<510>

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cgatatatta caggogteaa teacacatee gagtaaaaag ttattgtetg ttgaatttge 360

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| የ ታታ | | | | | 388 DNA | <210> <211> <212> |
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<510>

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अर्ट्याचित व्यव्याचित्वाचे व्यव्यवाचित्वे व्यव्यवाच्या र्ट्याच्यव्य र्ट्याच्यव्य र्ट्याच्यव्य

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| cattaattgt | ttttctttac | ςτεςεςτες | attgttgtt | CLLCALLLL | ctccatgtat | 150 |
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| гатсадсаса | ccrdcscccc | მმმყբբმმმբ | ttatgccttt | агдагдсгда | | 0TÞ |
| cdcrddrcdc | ςρετταςςςς | ραδοροστρο | ааатдтааса | rrdsrdrsdc | дгггэгэдсэ | 360 |
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| дгагдгдсяя | травтддстс | сасасааааt | сгааадгссд | craacgagca | taatcatcct | 180 |
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| ನೆನಿಕ್ಕೆನಿತತಾರ್ಧ | τοοταοτττ | attcgatgac | сдгэээдгдд | гасстдгада | гатдгассдд | 09 |
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| ςαςαςσαςας | ςααςααςς | астдатстас | attgttgatt | attaacagtg | actgccacat | 09 |
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 ${\tt Gl} {\tt Xcine} \ {\tt max}$

20280 403 DNA <213>

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<510> 20285 arrac 182 fictaaccag caagicotic agiigaaga taticaggot aagotaigg aataggotaa 180 150 taaggittaag cagcigcagc aaaagcigaa igaatiggaa gaticcaigc cicaccaacc гааддагсгс авасваядс гдагаддсгд дадсададаг асгдгаддгд асггдсдссгг 09 2028₫ <007> Glycine max <213> DNA <212> **T82** <5112> 2028₫ <210> tttctcctc ttttcctct acgacatgtc gaacaatact 09ħ rararradas sestecest esetadas cestracas costeses restagace **450** tattgacttg gatttttaat caacacctt ctcgtcccaa attcaaaatt gattcgcaaa 360 cfffraaat atagggfgtt aataactgca caagactatt gacggccaac acttctctta 300 daattactet caaatetgea eettteetag atgitaatta aeeegeegta aeaeeaett .5**₹**0 эвысстсява тупите вупительного поста под поста под поста под поста под поста п 180 dedercorer esadrogaco raredoser fredorffo recreaçada didicipoco 150 09 20283 <007> unsure at all n locations <222> Glycine max <213> ANG <212> <211> 09₺ 20283 <510> agteatgtet agteatgaea etegteaeat 057 drogasasa accteaaaa tetagadiga etegticaee tatigtatit etgicaaaga **450** tridictide aatiliatig gergaacet aaaceataa atictiacaa aaacattaaa 360

teaatatet tagtettgtt ettgaaceat gaattgtget gagtitaagt teetttgagt

attetttaga atttaeaeeg attaaaettg etatagaage tagatttgae tttetatgu

300

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| 300 | crcsrdrrrc | ccttttacca | τρασαρατρα | дсдсяясдся | ataggcttt | гдсгсгсяяг |
| 240 | гсдгдггчч | ddragtacac | actettagät | cscgtgttta | acactctact | ddsccrcscs |
| 180 | гааддсстса | таааааттат | ааасусаа | acaggaaaa | ccataattca | ccrdcrccsd |
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| 22.5 | | • | | | | ೧೯೦ಡಿಡಿಇಡಿಂ |
| 898 | | | | | | |
| 390 | gregaatttg | αρασρρασο | cgagttagaa | atcagacatt | acgggactca | tegatetatt |
| 300 | гасаадсдъд | аасаттсаат | гдздздсггэ | rrdsatttgc | rrarrgregr | адгаааааа |
| 5₫0 | catacatctg | dddactcaat | . פפרפרפררכ | cgatcgtctc | ааттсаатат | адаасттсат |
| 180 | дааттатстс | attgtcgctt | асадааадсь | agacatccga | ддясрсяярс | atatattacg |
| 150 | дядсарсряд | attcaatttc | døddcccøcø | аассадсссс | τοττοαδττα | taaaatgtta |
| 09 | dscstccdsd | gactcaatca | tatattacgg | адсдісісаа | tragatttag | адсгрсадга |
| | | , | | | 20285 | <004> |
| | | | | 3 | DNA Glycine max | <212> |
| | | | | | | |

| 300 | адававсста | свасудаваа | ссдсассава | tacsagtatt | ε α εεεεααε | cgcatcttca |
|-----|------------|------------|-------------|--------------|-----------------------------|-------------------------|
| 5₫0 | ссааасатст | гддгаддааа | аастаттсаа | actccatatg | tttaaagaaa | гаадасссат |
| 180 | адддсгргая | татсатдаат | стдсятдста | дссдагаага | градасгач | свавататт |
| 750 | actectette | аставатда | сғадаассғ | даадааассд | τασατοτττα | 222323333 |
| 09 | ddsdrrdrrd | ссаасаааат | cragaacccc | ttttctgcac | гдвасатаас | адсградсыг |
| | | | | | 20289 | <004> |
| | | | | > | 20289 DNA Glycine max | <211> <211> <213> |
| τsħ | | | đ | псссастаа | teattteat | ггдээдггдд |
| 420 | ааатаддстп | гсдгссясяя | сгадісаадс | atcaactcgc | гадддсгааа | ааадтдааас |
| 360 | attaataacg | аассдаассд | teteattett | crraddarr | dacgttttct | ааатсаатсд |
| 300 | агааадсдда | ттттадсява | адававаевес | ataataataa | авававатавт | аааадаадс |
| 5₫0 | гддваатсаа | draaccacgr | cddrcdrscc | гссдяссдгг | таватават | rascreedge |
| 180 | tattatccgt | дсссдвассд | рссвссдвас | аваатаватт | асттавават | ttctcgctta |
| 150 | cttaagtcat | адссаттта | аасдідсііа | ςςςςςςτςτ | acttttata | כמברברמברב |
| 09 | בבבבפבבכב | ατττοαταος | tecacattgt | cccdragtgg | attctatgta | ctttcgattc |
| | | | suoj | all n locati | SOS88 nuente et e | <2223> <400> |
| | | | | > | Clycine may | <211> <213> |
| 393 | | | đđ¢ | tttgcaagtt | atatttggcc | аааддсьдса |
| 360 | attcatgctc | гаадатассд | atattgccaa | attcatcatg | сттсстватс | ddccrccsat |
| 300 | gatgagaatt | tcttattgat | cagctggtga | ttttgttat | саатаасата | εεεδεοάδαε |
| 240 | tctctatatt | стссаттдав | ассаадсдсс | tgcctcttca | gaccattcca | гдзагдагда |
| 180 | tttctagagg | ataggaggtt | гааааадаад | агггасавва | בבמבבבבמ | בפרבפרפר |

<211> 444

| 868 | | | בבכ | ттссаваста | тсаааастт | асдсасстас |
|-----|------------|------------------|------------|-------------|-----------------------------|-------------------------|
| 360 | аггадггса | ггадаасаад | ttttcaaaat | аааасдасас | дсссвассвс | actatacctt |
| 300 | агаассгэээ | tggaaataat | atacccttat | atctacttca | ttttgcagg | асавсадава |
| 240 | свассдева | ಕಡಿಡಿಡಿದ್ದಕತಡಿಡಿ | cdcscscsag | tcacacctct | ೦ಇತರಿತನಿತನಿತ | tctgcatgac |
| 180 | ссгдгэдээг | rraraaagcg | сгаааагссс | αασταστττ | scsdrddssd | aacaagttgg |
| ISO | crccrrrdc | caaactccac | tcattaaagt | tttgcatctg | caaaagcctc | ссдаддсаас |
| 09 | caacgetett | ವಿಡಿದ್ದಿಡಿದ್ದೇ | tagatgatag | σεςαααεαςς | αραρρασσ | адсссасавд |
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| ΤÞÞ | | | | a | тддттстсяя | attctataat |
| 420 | ttatttctaa | tatcattaat | aaccttacat | псасасадс | аасадсаасс | ааатдаатда |
| 360 | aataataaat | ссаадаааас | tttacacgga | attgttgtct | ссстапаата | actctttacg |
| 300 | аатастстда | сатдтаатас | ttgaccattg | ataagactcc | ttcataagat | асадссадас |
| 540 | aattcatagt | сггдсядггд | дсяддряяя | tettgtatga | dededcdedr | agacttcatt |
| 180 | crcsattggc | aacaatgatc | ddcsdsscdr | aacaataggt | αττττττα | aatctccctt |
| 150 | cttcttaagt | гвдсгвдгсг | αττττταστ | getttatt | taagttttgt | Lggaccaatc |
| 09 | гааддсгааа | ctatagttca | attattatac | адсаааасаа | ברבלברבר | псддаасдаа |
| | | | suoį | all n locat | 20290 | <223> |

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Glycine max

ANG

441 50590 <212>

<111>

<210>

ddsgagdrffg argcraadfa agardaagga arrrccffg greatcfff gcaragrasa 360

accetgasac aageteecga gggtgeetat cataattiga aaccettate aegeteed 180 cettaggggtgas ageteecga gggtgeetat cataattiga aaccettatg cteteteaga 120 cettaggggettat aaggaettat cataattiga aaccettatg cteteteaga 120

| ₹6202 | <004> |
|-------|-------|
| VOCUC | \UU\/ |

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ANG <SIS>

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<SIO> <SIO>

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<400>

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